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INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

| (51) International Patent Classification 6: | A2 | (11) International Publication Number: | WO 95/11984 |
|---|----|--|-----------------------|
| C12N 15/86, 7/01, 5/10, A61K 48/00, C07K 14/47 | | (43) International Publication Date: | 4 May 1995 (04.05.95) |

(21) International Application Number: PCT/US94/12235

(22) International Filing Date: 25 October 1994 (25.10.94)

(30) Priority Data:

08/142,669 25 October 1993 (25.10.93) US 08/246,007 19 May 1994 (19.05.94) US

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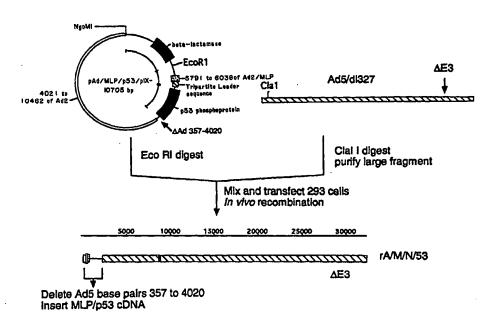
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Published

Without international search report and to be republished upon receipt of that report.

(54) Title: RECOMBINANT ADENOVIRAL VECTOR AND METHODS OF USE



(57) Abstract

This invention provides a recombinant adenovirus expression vector characterized by the partial or total deletion of the adenoviral protein IX DNA and having a gene encoding a foreign protein or a functional fragment or mutant thereof. Transformed host cells and a method of producing recombinant proteins and gene therapy also are included within the scope of this invention. Thus, for example, the adenoviral vector of this invention can contain a foreign gene for the expression of a protein effective in regulating the cell cycle, such as p53, Rb, or mitosin, or in inducing cell death, such as the conditional suicide gene thymidine kinase. (The latter must be used in conjunction with a thymidine kinase metabolite in order to be effective).

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RECOMBINANT ADENOVIRAL VECTOR AND METHODS OF USE BACKGROUND OF THE INVENTION

This application is a continuation-in-part of U.S. Serial No. 08/233,777, filed May 19, 1994, which is a continuation-in-part of U.S. Serial No. 08/142,669 filed October 25, 1993, the contents of which are hereby incorporated by reference into the present disclosure.

Throughout this application, various publications are referred to by citations within parentheses and in the bibliographic description, immediately preceding the claims. The disclosures of these publications are hereby incorporated by reference into the present disclosure to more fully describe the state of the art to which this invention pertains.

Production of recombinant adenoviruses useful for gene therapy requires the use of a cell line capable of supplying in trans the gene products of the viral E1 region which are deleted in these recombinant viruses. At present the only useful cell line available is the 293 cell line originally described by Graham et al. in 1977. 293 cells contain approximately the left hand 12% (4.3 kb) of the adenovirus type 5 genome (Aiello (1979) and Spector (1983)).

Adenoviral vectors currently being tested for 25 gene therapy applications typically are deleted for Ad2 or Ad5 DNA extending from approximately 400 base pairs from the 5' end of the viral genome to approximately 3.3 kb from the 5' end, for a total E1 deletion of 2.9 kb. Therefore, there exists a limited region of homology of approximately 1 kb between the DNA sequence of the recombinant virus and the Ad5 DNA within the cell line. This homology defines a region of potential recombination between the viral and cellular adenovirus sequences. Such a recombination results in a phenotypically wild-type virus bearing the Ad5 E1 region from the 293 cells. This recombination event

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presumably accounts for the frequent detection of wild-type adenovirus in preparations of recombinant virus and has been directly demonstrated to be the cause of wild-type contamination of the Ad2 based recombinant virus Ad2/CFTR-1 (Rich et al. (1993)).

Due to the high degree of sequence homology within the type C adenovirus subgroup such recombination is likely to occur if the vector is based on any group C adenovirus (types 1, 2, 5, 6).

In small scale production of recombinant adenoviruses, generation of contaminating wild-type virus can be managed by a screening process which discards those preparations of virus found to be contaminated. As the scale of virus production grows to meet expected demand for genetic therapeutics, the likelihood of any single lot being contaminated with a wild-type virus also will rise as well as the difficulty in providing non-contaminated recombinant preparations.

There will be over one million new cases of 20 cancer diagnosed this year, and half that number of cancerrelated deaths (American Cancer Society, 1993). mutations are the most common genetic alteration associated with human cancers, occurring in 50-60% of human cancers (Hollstein et al. (1991); Bartek et al. (1991); Levine 25 (1993)). The goal of gene therapy in treating p53 deficient tumors, for example, is to reinstate a normal, functional copy of the wild-type p53 gene so that control of cellular proliferation is restored. p53 plays a central role in cell cycle progression, arresting growth so that 30 repair or apoptisis can occur in response to DNA damage. Wild-type p53 has recently been identified as a necessary component for apoptosis induced by irradiation or treatment with some chemotherapeutic agents (Lowe et al. (1993) A and B). Due to the high prevalence of p53 mutations in human

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tumors, it is possible that tumors which have become refractory to chemotherapy and irradiation treatments may have become so due in part to the lack of wild-type p53. By resupplying functional p53 to these tumors, it is reasonable that they now are susceptible to apoptisis normally associated with the DNA damage induced by radiation and chemotherapy.

One of the critical points in successful human tumor suppressor gene therapy is the ability to affect a significant fraction of the cancer cells. The use of retroviral vectors has been largely explored for this purpose in a variety of tumor models. For example, for the treatment of hepatic malignancies, retroviral vectors have been employed with little success because these vectors are not able to achieve the high level of gene transfer required for in vivo gene therapy (Huber, B.E. et al., 1991; Caruso M. et al., 1993).

To achieve a more sustained source of virus production, researchers have attempted to overcome the problem associated with low level of gene transfer by direct injection of retroviral packaging cell lines into solid tumors (Caruso, M. et al., 1993; Ezzidine, Z.D. et al., 1991; Culver, K.W. et al., 1992). However, these methods are unsatisfactory for use in human patients 25 because the method is troublesome and induces inflammatory response against the packaging cell line in the patient. Another disadvantage of retroviral vectors is that they require dividing cells to efficiently integrate and express the recombinant gene of interest (Huber, B.E. 1991). Stable integration into an essential host gene can lead to the development or inheritance of pathogenic diseased states.

Recombinant adenoviruses have distinct advantages over retroviral and other gene delivery methods (for

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review, see Siegfried (1993)). Adenoviruses have never been shown to induce tumors in humans and have been safely used as live vaccines (Straus (1984)). Replication deficient recombinant adenoviruses can be produced by 5 replacing the E1 region necessary for replication with the target gene. Adenovirus does not integrate into the human genome as a normal consequence of infection, thereby greatly reducing the risk of insertional mutagenesis possible with retrovirus or adeno-associated viral (AAV) 10 vectors. This lack of stable integration also leads to an additional safety feature in that the transferred gene effect will be transient, as the extrachromosomal DNA will be gradually lost with continued division of normal cells. Stable, high titer recombinant adenovirus can be produced 15 at levels not achievable with retrovirus or AAV, allowing enough material to be produced to treat a large patient Moreover, adenovirus vectors are capable of population. highly efficient in vivo gene transfer into a broad range of tissue and tumor cell types. For example, others have 20 shown that adenovirus mediated gene delivery has a strong potential for gene therapy for diseases such as cystic fibrosis (Rosenfeld et al. (1992); Rich et al. (1993)) and α_1 -antitrypsin deficiency (Lemarchand et al. Although other alternatives for gene delivery, such as 25 cationic liposome/DNA complexes, are also currently being explored, none as yet appear as effective as adenovirus mediated gene delivery.

As with treating p53 deficient tumors, the goal of gene therapy for other tumors is to reinstate control of cellular proliferation. In the case of p53, introduction of a functional gene reinstates cell cycle control allowing for apoptotic cell death induced by therapeutic agents. Similarly, gene therapy is equally applicable to other tumor suppressor genes which can be used either alone or in combination with therapeutic agents to control cell cycle progression of tumor cells and/or induce cell death.

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Moreover, genes which do not encode cell cycle regulatory proteins, but directly induce cell death such as suicide genes or, genes which are directly toxic to the cell can be used in gene therapy protocols to directly eliminate the cell cycle progression of tumor cells.

Regardless of which gene is used to reinstate the control of cell cycle progression, the rationale and practical applicability of this approach is identical. Namely, to achieve high efficiencies of gene transfer to express therapeutic quantities of the recombinant product. The choice of which vector to use to enable high efficiency gene transfer with minimal risk to the patient is therefore important to the level of success of the gene therapy treatment.

Thus, there exists a need for vectors and methods which provide high level gene transfer efficiencies and protein expression which provide safe and effective gene therapy treatments. The present invention satisfies this need and provides related advantages as well.

20 <u>SUMMARY OF THE INVENTION</u>

This invention provides a recombinant adenovirus expression vector characterized by the partial or total deletion of the adenoviral protein IX DNA and having a gene encoding a foreign protein or a functional fragment or mutant thereof. Transformed host cells and a method of producing recombinant proteins and gene therapy also are included within the scope of this invention.

Thus, for example, the adenoviral vector of this invention can contain a foreign gene for the expression of a protein effective in regulating the cell cycle, such as p53, Rb, or mitosin, or in inducing cell death, such as the conditional suicide gene thymidine kinase. (The latter

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must be used in conjunction with a thymidine kinase metabolite in order to be effective).

BRIEF DESCRIPTION OF THE FIGURES

Figure 1 shows a recombinant adenoviral vector of this invention. This construct was assembled as shown in Figure 1. The resultant virus bears a 5' deletion of adenoviral sequences extending from nucleotide 356 to 4020 and eliminates the Ela and Elb genes as well as the entire protein IX coding sequence, leaving the polyadenylation site shared by the Elb and pIX genes intact for use in terminating transcription of any desired gene.

Figure 2 shows the amino acid sequence of p110RB.

Figure 3 shows a DNA sequence encoding a retinoblastoma tumor suppressor protein.

15 schematic Figure 4 shows of recombinant p53/adenovirus constructs within the scope of invention. The p53 recombinants are based on Ad 5 and have had the El region of nucleotides 360-3325 replaced with a 1.4 kb full length p53 cDNA driven by the Ad 2 MLP (A/M/53)20 or human CMV (A/C/53) promoters followed by the Ad 2 tripartite leader cDNA. The control virus A/M has the same Ad 5 deletions as the A/M/53 virus but lacks the 1.4 kb p53 cDNA insert. The remaining Elb sequence (705 nucleotides) have been deleted to create the protein IX deleted 25 constructs A/M/N/53 and A/C/N/53. These constructs also have a 1.9 kb Xba I deletion within adenovirus type 5 region E3.

Figures 5A and 5B show p53 protein expression in tumor cells infected with A/M/53 and A/C/53. Figure 5A)

30 Saos-2 (osteosarcoma) cells were infected at the indicated multiplicities of infection (MOI) with either the A/M/53 or

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A/C/53 purified virus and harvested 24 hours later. The p53 antibody pAb 1801 was used to stain immunoblots of samples loaded at equal total protein concentrations. Equal protein concentration of SW480 cell extracts, which overexpress mutant p53 protein, were used as a marker for p53 size. "O" under the A/C/53 heading indicates a mock infection, containing untreated Saos-2 lysate. Figure 5B) Hep 3B (hepatocellular carcinoma) cells were infected with the A/M/53 or A/C/53 virus at the indicated MOI and analyzed as in part A.) The arrow indicates the position of the p53 protein.

Figures 6A through 6C show p53 dependent Saos-2 morphology change. Subconfluent (1 x 10⁵ cells/10 cm plate) Saos-2 cells were either uninfected (A), infected at an MOI = 50 with (B) the control A/M virus or (C) the A/C/53 virus. The cells were photographed 72 hours post-infection.

Figure 7 shows p53 dependent inhibition of DNA synthesis in human tumor cell lines by A/M/N/53 and A/C/N/53. Nine different tumor cell lines were infected with either control adenovirus A/M (-x-x-), or the p53 expressing A/M/N/53 (-Δ-Δ-), or A/C/N/53 (-O-O-) virus at increasing MOI as indicated. The tumor type and p53 status is noted for each cell line (wt = wild type, null = no protein expressed, mut = mutant protein expressed). DNA synthesis was measured 72 hours post-infection as described below in Experiment No. II. Results are from triplicate measurements at each dose (mean+/- SD), and are plotted as % of media control versus MOI. * H69 cells were only tested with A/M and A/M/N/53 virus.

Figure 8 shows tumorigenicity of p53 infected Saos-2 cells in nude mice. Saos-2 cells were infected with either the control A/M virus or the p53 recombinant A/M/N/53 at MOI = 30. Treated cells were injected

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subcutaneously into the flanks of nude mice, and tumor dimensions were measured (as described in Experiment No. II) twice per week for 8 weeks. Results are plotted as tumor size versus days post tumor cell implantation for both control A/M (-x-x-) and A/M/N/53 (-Δ-Δ-) treated cells. Error bars represent the mean tumor size =/- SEM for each group of 4 animals at each time point.

Figure 9 is expression of rAd/p53 H69 (SCLC) cells were injected established tumors. 10 subcutaneously into nude mice and allowed to develop tumors for 32 days until reaching a size of approximately 25-50 mm3. Mice were randomized and injected peritumorally with 2 x 10° pfu of either control A/C/S-gal or A/C/53 virus. Tumors were excised 2 and 7 days post injection, and polyA 15 RNA was prepared from each tumor sample. RT-PCR was carried out using equal RNA concentrations and primers specific for recombinant p53 message. PCR amplification was for 30 cycles at 94°C 1 min., 55°C 1.5 min., 72°C 2 min., and a 10 min., 72°C final extension period in an Omnigen 20 thermalcycler (Hybaid). The PCR primers used were a 5' Tripartite Leader cDNA (5' - CGCCACCGAGGGACCTGAGCGAGTC-3') and a 3' p53 primer (5' - TTCTGGGAAGGGACAGAAGA-3'). Lanes 1, 2, 4, and 5 are p53 treated samples excised at day 2 or Lanes 3 and 6 are from ß-gal treated 7 as indicated. 25 tumors. Lanes 7,8, and 9 are replicates of lanes 4,5, and 6 respectively, amplified with actin primers to verify equal loading. Lane 10 is a positive control using a tripartite/p53 containing plasmid.

Figures 10A and 10B show in vivo tumor suppression and increased survival time with A/M/N/53. H69 (SCLC) tumor cells were injected subcutaneously into nude mice and allowed to develop for 2 weeks. Peritumoral injections of either buffer alone (---), control A/M adenovirus (-x-x-), or A/M/N/53 (-Δ-Δ), both viruses (2 x 10° pfu/injection) were administered twice per week for a

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total of 8 doses. Tumor dimensions were measured twice per week and tumor volume was estimated as described in Experiment No. II. A) Tumor size is plotted for each virus versus time (days) post inoculation of H69 cells. Error bars indicate the mean tumor size +/- SEM for each group of 5 animals. Arrows indicate days virus injections. B) Mice were monitored for survival and the fraction of mice surviving per group versus time post inoculation of buffer alone (----), control A/M (··· ···) or A/M/N/53 (—) virus treated H69 cells is plotted.

Figures 11A through 11C show maps of recombinant plasmid constructions. Plasmids were constructed as detailed in below. Bold lines in the constructs indicate genes of interest while boldface type indicates the 15 restriction sites used to generate the fragments to be ligated together to form the subsequent plasmid indicated by the arrows. In Figure 11A, the plasmid pACNTK was constructed by subcloning the HSV-TK gene from pMLBKTK (ATCC No. 39369) into the polylinker of a cloning vector, 20 followed by isolation of the TK gene with the desired ends for cloning into the pACN vector. The pACN vector contains adenoviral sequences necessary for in vivo recombination to occur to form recombinant adenovirus (see Figure 12). Figure 11B, the construction of the plasmid pAANTK is shown 25 beginning with PCR amplified fragments encoding the α fetoprotein enhancer (AFP-E) and promoter (AFP-P) regions subcloned through several steps into a final plasmid where the AFP enhancer and promoter are upstream of the HSV-TK gene followed by adenovirus Type 2 sequences necessary for in vivo recombination to occur to form recombinant 30 adenovirus. In Figure 11C, the construction of the plasmid pAANCAT is shown beginning with the isolation of the chloramphenicol acetyltransferase (CAT) gene from a commercially available plasmid and subcloning it into the pAAN plasmid (see above), generating the final plasmid PAANCAT enhancer/promoter where the AFP

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transcription of the CAT gene in an adenovirus sequence background.

Figure 12 is a schematic map of recombinant adenoviruses ACNTK, AANTK and AANCAT. To construct 5 recombinant adenoviruses from the plasmids described in Figure 11, 4 parts (20 μ g) of either plasmid pACNTK, pAANTK, or pAANCAT were linearized with Eco R1 cotransfected with 1 part (5 μ g) of the large fragment of Cla 1 digested recombinant adenovirus (rACS-gal) containing 10 an E3 region deletion (Wills et al., 1994). resulting viruses, the Ad 5 nucleotides 360 - 4021 are replaced by either the CMV promoter and tripartite leader cDNA (TPL) or the α -fetoprotein enhancer and promoter (AFP) driving expression of the HSV-1 TK or CAT gene as 15 indicated. The resulting recombinant adenoviruses are designated ACNTK, AANTK, and AANCAT respectively.

Figure 13 shows promoter specificity of CAT expression in the recombinant adenoviral vectors. Two (2) X 106 of the designated cell lines were infected at MOIs = 30 or 100 of the recombinant adenovirus AANCAT as indicated or left uninfected (UN). Hep G2 and Hep 3B cells express α -fetoprotein whereas the other cell lines do not. After three days, the cells were harvested, extract volumes were adjusted for equal total protein concentrations, and CAT 25 activity was measured as described in Methods section. An equal number of uninfected cells served as individual controls for background CAT activity, while 14C labelled chloramphenicol (14C-only) and extract from a stable cell line (B21) expressing CAT activity served as 30 negative and positive controls respectively. conversion of acetyl CoA is indicated, demonstrating that CAT expression is limited to those cells expressing α fetoprotein.

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Figure 14 shows the effects of TK/GCV treatment on hepatocellular carcinoma cell lines and the effects of promoter specificity. Hep-G2 (AFP positive) and HLF (AFP negative) cell lines were infected overnight with ACNTK [-5 Δ-] AANTK [-▲-], or control ACN [-□-] virus at an infection multiplicity of 30 and subsequently treated with a single dose of ganciclovir at the indicated concentrations. Cell proliferation was assessed by adding 3H-thymidine to the cells approximately 18 hours prior to harvest. 3H-thymidine 10 incorporation into cellular nucleic acid was measured 72 hours after infection (Top Count, Packard and expressed as a percent (mean +/- S.D.) of untreated control. results show a non-selective dose dependent inhibition of proliferation with the CMV driven construct, while AFP 15 driven TK selectively inhibits Hep-G2.

Figure 15 shows cytotoxicity of ACNTK plus ganciclovir in HCC. HLF cells were infected at an MOI of 30 with either ACNTK [-•-] or the control virus ACN [-□-] and treated with ganciclovir at the indicated doses.

20 Seventy-two (72) hours after ganciclovir treatment, the amount of lactate dehydrogenase (LDH) released into the cell supernatant were measured colorimetrically and plotted (mean+/-SEM) versus ganciclovir concentration for the two virus treated groups.

Figures 16A and 16B show the effect of ACNTK plus ganciclovir on established hepatocellular carcinoma (HCC) tumors in nude mice. One (1) X 10⁷ Hep 3B cells were injected subcutaneously into the flank of female nude mice and allowed to grow for 27 days. Mice then received intratumoral and peritumoral injections of either the ACNTK [-Φ-] or control ACN [-□-] virus (1 X 10⁹ iu in 100 μl volume) every other day for a total of three doses (indicated by arrows). Injections of ganciclovir (100 mg/kg ip) began 24 hours after the initial virus dose and continued for a total of 10 days. In Figure 6A, tumor

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sizes are plotted for each virus versus days post infection (mean +/- SEM). In Figure 6B, body weight for each virustreated animal group is plotted as the mean +/-SEM versus days post infection.

DETAILED DESCRIPTION OF THE INVENTION

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To reduce the frequency of contamination with wild-type adenovirus, it is desirable to improve either the virus or the cell line to reduce the probability of For example, an adenovirus from a group recombination. 10 with low homology to the group C viruses could be used to engineer recombinant viruses with little propensity for recombination with the Ad5 sequences in 293 cells. However, an alternative, easier means of reducing the recombination between viral and cellular sequences is to 15 increase the size of the deletion in the recombinant virus and thereby reduce the extent of shared sequence between it and the Ad5 genes in the 293 cells.

Deletions which extend past 3.5 kb from the 5' end of the adenoviral genome affect the gene for adenoviral 20 protein IX and have not been considered desirable in adenoviral vectors (see below).

The protein IX gene of the adenoviruses encodes a minor component of the outer adenoviral capsid which stabilizes the group-of-nine hexons which compose the 25 majority of the viral capsid (Stewart (1993)). Based upon study of adenovirus deletion mutants, protein IX initially was thought to be a non-essential component of the adenovirus, although its absence was associated with greater heat lability than observed with wild-type virus (Colby and Shenk (1981)). More recently it was discovered that protein IX is essential for packaging full length viral DNA into capsids and that in the absence of protein IX, only genomes at least 1 kb smaller than wild-type could

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be propagated as recombinant viruses (Ghosh-Choudhury et al. (1987)). Given this packaging limitation, protein IX deletions deliberately have not been considered in the design of adenoviral vectors.

In this application, reference is made to standard textbooks of molecular biology that contain definitions, methods and means for carrying out basic techniques, encompassed by the present invention. See for example, Sambrook et al. (1989) and the various references cited therein. This reference and the cited publications are expressly incorporated by reference into this disclosure.

Contrary to what has been known in the art, this invention claims the use of recombinant adenoviruses 15 bearing deletions of the protein IX gene as a means of reducing the risk of wild-type adenovirus contamination in virus preparations for use in diagnostic and therapeutic applications such as gene therapy. As used herein, the term "recombinant" is intended to mean a progeny formed as 20 the result of genetic engineering. These deletions can remove an additional 500 to 700 base pairs of DNA sequence that is present in conventional E1 deleted viruses (smaller, less desirable, deletions of portions of the pIX gene are possible and are included within the scope of this 25 invention) and is available for recombination with the Ad5 sequences integrated in 293 cells. Recombinant adenoviruses based on any group C virus, serotype 1, 2, 5 and 6, are included in this invention. Also encompassed by this invention is a hybrid Ad2/Ad5 based recombinant virus expressing the human p53 cDNA from the adenovirus type 2 major late promoter. This construct was assembled as shown in Figure 1. The resultant virus bears a 5' deletion of adenoviral sequences extending from about nucleotide 357 to 4020 and eliminates the Ela and Elb genes as well as the entire protein IX coding sequence, leaving

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polyadenylation site shared by the Elb and protein IX genes intact for use in terminating transcription of any desired A separate embodiment is shown in Figure 4. Alternatively, the deletion can be extended an additional 5 30 to 40 base pairs without affecting the adjacent gene for protein IVa2, although in that case an exogenous polyadenylation signal is provided to terminate transcription of genes inserted into the recombinant virus. The initial virus constructed with this deletion is easily 10 propagated in 293 cells with no evidence of wild-type viral contamination and directs robust p53 expression from the transcriptional unit inserted at the site of the deletion.

The insert capacity of recombinant viruses bearing the protein IX deletion described above is 15 approximately 2.6 kb. This is sufficient for many genes including the p53 cDNA. Insert capacity can be increased introducing other deletions into the adenoviral backbone, for example, deletions within early regions 3 or 4 (for review see: Graham and Prevec (1991)). example, the use of an adenoviral backbone containing a 1.9 kb deletion of non-essential sequence within early region 3. With this additional deletion, the insert capacity of the vector is increased to approximately 4.5 kb, large enough for many larger cDNAs, including that of the 25 retinoblastoma tumor suppressor gene.

recombinant adenovirus expression vector characterized by the partial or total deletion of the adenoviral protein IX DNA and having a gene encoding a foreign protein, or a functional fragment or mutant thereof is provided by this invention. These vectors are useful for the safe recombinant production of diagnostic and therapeutic polypeptides and proteins, and importantly, for the introduction of genes in gene therapy. Thus, for example, the adenoviral vector of this invention 35 can contain a foreign gene for the expression of a protein

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effective in regulating the cell cycle, such as p53, Rb, or mitosin, or in inducing cell death, such as the conditional suicide gene thymidine kinase. (The latter must be used in conjunction with a thymidine kinase metabolite in order to 5 be effective). Any expression cassette can be used in the vectors of this invention. An "expression cassette" means a DNA molecule having a transcription promoter/enhancer such as the CMV promotor enhancer, etc., a foreign gene, and in some embodiments defined below, a polyadentlyation As used herein, the term "foreign gene" is 10 signal. intended to mean a DNA molecule not present in the exact orientation and position as the counterpart DNA molecule found in wild-type adenovirus. The foreign gene is a DNA molecule up to 4.5 kilobases. "Expression vector" means a 15 vector that results in the expression of inserted DNA sequences when propagated in a suitable host cell, i.e., the protein or polypeptide coded for by the DNA is synthesized by the host's system. The recombinant adenovirus expression vector can contain part of the gene 20 encoding adenovirus protein IX, provided that biologically active protein IX or fragment thereof is not produced. Example of this vector are an expression vector having the restriction enzyme map of Figures 1 or 4.

Inducible promoters also can be used in the 25 adenoviral vector of this invention. These promoters will initiate transcription only in the presence of additional molecule. Examples of inducible promoters include those obtainable from a ß-interferon gene, a heat shock gene, a metallothionine gene or those obtainable from steroid hormone-responsive genes. Tissue specific expression has been well characterized in the field of gene expression and tissue specific and inducible promoters such as these are very well known in the art. These genes are used to regulate the expression of the foreign gene after it has been introduced into the target cell.

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Also provided by this invention is a recombinant adenovirus expression vector, as described above, having less extensive deletions of the protein IX gene sequence extending from 3500 bp from the 5' viral termini to approximately 4000 bp, in one embodiment. In a separate embodiment, the recombinant adenovirus expression vector can have a further deletion of a non-essential DNA sequence in adenovirus early region 3 and/or 4 and/or deletion of the DNA sequences designated adenovirus E1a and E1b. In this embodiment, foreign gene is a DNA molecule of a size up to 4.5 kilobases.

A further embodiment has a deletion of up to forty nucleotides positioned 3' to the Ela and Elb deletion and pIX and a foreign DNA molecule encoding a polyadenylation signal inserted into the recombinant vector in a position relative to the foreign gene to regulate the expression of the foreign gene.

For the purposes of this invention, the recombinant adenovirus expression vector can be derived 20 from wild-type group adenovirus, serotype 1, 2, 5 or 6.

In one embodiment, the recombinant adenovirus expression vector has a foreign gene coding for a functional tumor suppressor protein, or a biologically active fragment thereof. As used herein, the term "functional" as it relates to a tumor suppressor gene, refers to tumor suppressor genes that encode tumor suppressor proteins that effectively inhibit a cell from behaving as a tumor cell. Functional genes can include, for instance, wild type of normal genes and modifications of normal genes that retains its ability to encode effective tumor suppressor proteins and other anti-tumor genes such as a conditional suicide protein or a toxin.

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Similarly, "non-functional" as used herein is synonymous with "inactivated." Non-functional or defective genes can be caused by a variety of events, including for example point mutations, deletions, methylation and others known to those skilled in the art.

As used herein, an "active fragment" of a gene includes smaller portions of the gene that retain the ability to encode proteins having tumor suppressing activity. p56^{RB}, described more fully below, is but one example of an active fragment of a functional tumor suppressor gene. Modifications of tumor suppressor genes are also contemplated within the meaning of an active fragment, such as additions, deletions or substitutions, as long as the functional activity of the unmodified gene is retained.

Another example of a tumor suppressor gene is retinoblastoma (RB). The complete RB cDNA nucleotide sequences and predicted amino acid sequences of the resulting RB protein (designated p110RB) are shown in Lee et 20 al. (1987) and in Figure 3. Also useful to express retinoblastoma tumor suppressor protein is a DNA molecule encoding the amino acid sequence shown in Figure 2 or having the DNA sequence shown in Figure 3. A truncated version of p110^{RB}, called p56^{RB} also is useful. sequence of p56^{RB}, see Huang et al. (1991). Additional tumor suppressor genes can be used in the vectors of this invention. For illustration purposes only, these can be pl6 protein (Kamb et al. (1994)), p21 protein, Wilm's tumor WT1 protein, mitosin, h-NUC, or colon carcinoma DCC 30 protein. Mitosin is described in X. Zhu and W-H Lee, U.S. Application Serial No. 08/141,239, filed October 22, 1993, and a subsequent continuation-in-part by the inventors, attorney docket number P-CJ 1191, filed October 1994, both of which are herein incorporated by 35 reference. Similarly, h-NUC is described by W-H Lee and P-

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L Chen, U.S. Application Serial No. 08/170,586, filed December 20, 1993, herein incorporated by reference.

As is known to those of skill in the art, the term "protein" means a linear polymer of amino acids joined 5 in a specific sequence by peptide bonds. As used herein. the term "amino acid" refers to either the D or L stereoisomer form of the amino acid, unless otherwise specifically designated. Also encompassed within the scope of this invention are equivalent proteins or equivalent 10 peptides, e.g., having the biological activity of purified wild type tumor suppressor protein. "Equivalent proteins" and "equivalent polypeptides" refer to compounds that depart from the linear sequence of the naturally occurring proteins or polypeptides, but which have amino acid substitutions that do not change its biologically activity. These equivalents can differ from the native sequences by the replacement of one or more amino acids with related amino acids, for example, similarly charged amino acids, or the substitution or modification of side chains or 20 functional groups.

Also encompassed within the definition of a functional tumor suppressor protein is any protein whose presence reduces the tumorigenicity, malignancy hyperproliferative phenotype of the host cell. Examples of 25 tumor suppressor proteins within this definition include, but are not limited to pl10RB, p56RB, mitosin, h-NUC and p53. "Tumorigenicity" is intended to mean having the ability to form tumors or capable of causing tumor formation and is synonymous with neoplastic growth. "Malignancy" 30 intended to describe a tumorigenic cell having the ability to metastasize and endanger the life of the host organism. "Hyperproliferative phenotype" is intended to describe a cell growing and dividing at a rate beyond the normal limitations of growth for that cell type. "Neoplastic" 35 also is intended to include cells lacking endogenous

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functional tumor suppressor protein or the inability of the cell to express endogenous nucleic acid encoding a functional tumor suppressor protein.

An example of a vector of this invention is a recombinant adenovirus expression vector having a foreign gene coding for p53 protein or an active fragment thereof is provided by this invention. The coding sequence of the p53 gene is set forth below in Table I.

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TABLE 1

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V*SHR PGSR* LLGSG DTLRS GWERA FHDGD TLPWI GSQTA FRVTA MEEPQ

- 5 SDPSV EPPLS QETFS DLWKL LPENN VLSPL PSQAM DDLML SPDDI EQWFT 150
 - EDPGP DEAPR MPEAA PPVAP APAAP TPAAP APAPS WPLSS SVPSQ KTYQG
 - SYGFR LGFLH SGTAK SVTCT YSPAL NKMFC QLAKT CPVQL WVDST PPPGT
 250
 - RVRAM AIYKQ SQHMT EVVRR CPHHE RCSDS DGLAP PQHLI RVEGN LRVEY
 - LDDRN TFRHS VVVPY EPPEV GSDCT TIHYN YMCNS SCMGG MNRRP ILTII
 350
- 15 TLEDS SGNLL GRNSF EVRVC ACPGR DRRTE EENLR KKGEP HHELP PGSTK
 400
 - RALPN NTSSS PQPKK KPLDG EYFTL QIRGR ERFEM FRELN EALEL KDAOA

GKEPG GSRAH SSHLK SKKGQ STSRH KKLMF KTEGP DSD*

* = Stop codon

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20 Any of the expression vectors described herein are useful as compositions for diagnosis or therapy. vectors can be used for screening which of many tumor suppressor genes would be useful in gene therapy. example, a sample of cells suspected of being neoplastic can be removed from a subject and mammal. The cells can then be contacted, under suitable conditions and with an effective amount of a recombinant vector of this invention having inserted therein a foreign gene encoding one of several functional tumor suppressor genes. Whether the 30 introduction of this gene will reverse the malignant phenotype can be measured by colony formation in soft agar or tumor formation in nude mice. If the malignant phenotype is reversed, then that foreign gene is determined to be a positive candidate for successful gene therapy for 35 the subject or mammal. When used pharmaceutically, they

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can be combined with one or more pharmaceutically acceptable carriers. Pharmaceutically acceptable carriers are well known in the art and include aqueous solutions such as physiologically buffered saline or other solvents or vehicles such as glycols, glycerol, vegetable oils (eg., olive oil) or injectable organic esters. A pharmaceutically acceptable carrier can be used to administer the instant compositions to a cell in vitro or to a subject in vivo.

A pharmaceutically acceptable carrier can contain 10 a physiologically acceptable compound that acts, example, to stabilize the composition or to increase or decrease the absorption of the agent. A physiologically acceptable compound can include, for example. 15 carbohydrates, such as glucose, sucrose or dextrans, antioxidants, such as ascorbic acid or glutathione, chelating agents, low molecular weight proteins or other stabilizers excipients. or Other physiologically acceptable compounds include wetting agents, emulsifying agents, dispersing agents or preservatives, which are 20 particularly useful for preventing the growth or action of microorganisms. Various preservatives are well known and include, for example, phenol and ascorbic acid. skilled in the art would know that the choice of a 25 pharmaceutically acceptable carrier, including physiologically acceptable compound, depends, for example, on the route of administration of the polypeptide and on the particular physio-chemical characteristics of the specific polypeptide. For example, a physiologically acceptable compound such as aluminum monosterate or gelatin is particularly useful as a delaying agent, which prolongs the rate of absorption of a pharmaceutical composition administered to a subject. Further examples of carriers, stabilizers or adjutants can be found in Remington's Pharm. Sci., 15th Ed. (Mack Publ. Co., Easton, 1975), incorporated herein by reference. The

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pharmaceutical composition also can be incorporated, if desired, into liposomes, microspheres or other polymer matrices (Gregoriadis, Liposome Technology, Vol. 1 (CRC Press, Boca Raton, Florida 1984), which is incorporated 5 herein by reference). Liposomes, for example, which consist of phospholipids or other lipids, are nontoxic, physiologically acceptable and metabolizable carriers that are relatively simple to make and administer.

As used herein, "pharmaceutical composition" 10 refers to any of the compositions of matte described herein combination with one ormore pharmaceutically acceptable carriers. The compositions can then be administered therapeutically or prophylactically. They can be contacted with the host cell in vivo, ex vivo, 15 or in vitro, in an effective amount. In vitro and ex vivo means of contacting host cells are provided below. of practiced in vivo. methods administering pharmaceutical containing the vector of this invention, are well known in the art and include but are not limited to, 20 administration orally, intra-tumorally, intravenously, intramuscularly or intraperitoneal. Administration can be effected continuously or intermittently and will vary with the subject and the condition to be treated, e.g., as is the case with other therapeutic compositions (Landmann et 25 al. (1992); Aulitzky et al. (1991); Lantz et al. (1990); Supersaxo et al. (1988); Demetri et al. (1989); and LeMaistre et al. (1991)).

Further provided by this invention transformed procaryotic or eucaryotic host cell, for 30 example an animal cell or mammalian cell, having inserted a recombinant adenovirus expression vector described above. Suitable procaryotic cells include but are not limited to bacterial cells such as E. coli cells. transforming host cells with retroviral vectors are known in the art, see Sambrook et al. (1989) and include, but are

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not limited to transfection, electroporation, and microinjection.

As used throughout this application, the term animal is intended to be synonymous with mammal and is to include, but not be limited to bovine, porcine, feline, simian, canine, equine, murine, rat or human. Additional host cells include but are not limited to any neoplastic or tumor cell, such as osteosarcoma, ovarian carcinoma, breast carcinoma, melanoma, hepatocarcinoma, lung cancer, brain cancer, colorectal cancer, hematopoietic cell, prostate cancer, cervical carcinoma, retinoblastoma, esophageal carcinoma, bladder cancer, neuroblastoma, or renal cancer.

Additionally, any eucaryotic cell line capable of expressing Ela and Elb or Ela, Elb and pIX is a suitable host for this vector. In one embodiment, a suitable eucaryotic host cell is the 293 cell line available from the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland, U.S.A. 20231.

Any of the transformed host cells described 20 herein are useful as compositions for diagnosis or therapy. When used pharmaceutically, they can be combined with various pharmaceutically acceptable carriers. Suitable pharmaceutically acceptable carriers are well known to those of skill in the art and, for example, are described above. The compositions can then be administered therapeutically or prophylactically, in effective amounts, described in more detail below.

A method of transforming a host cell also is provided by this invention. This method provides contacting a host cell, i.e., a procaryotic or eucaryotic host cell, with any of the expression vectors described herein and under suitable conditions. Host cells transformed by this method also are claimed within the

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scope of this invention. The contacting can be effected in vitro, in vivo, or ex vivo, using methods well known in the art (Sambrook et al. (1989)) and using effective amounts of the expression vectors. Also provided in this invention is 5 a method of producing a recombinant protein or polypeptide by growing the transformed host cell under suitable conditions favoring the transcription and translation of Methods of recombinant inserted foreign gene. expression in a variety of host cells, such as mammalian, 10 yeast, insect or bacterial cells, are widely known, including those described in Sambrook et al., supra. translated foreign gene can then be isolated by convention means, such as column purification or purification using an anti-protein antibody. The isolated protein or polypeptide 15 also is intended within the scope of this invention. used herein, purified or isolated mean substantially free of native proteins or nucleic acids normally associated with the protein or polypeptide in the native or host cell environment.

Also provided by this invention are non-human animals having inserted therein the expression vectors or transformed host cells of this invention. These "transgenic" animals are made using methods well known to those of skill in the art, for example as described in U.S.

25 Patent No. 5,175,384 or by conventional ex vivo therapy techniques, as described in Culver et al. (1991).

As shown in detail below, the recombinant adenoviruses expressing a tumor suppressor wild-type p53, as described above, can efficiently inhibit DNA synthesis and suppress the growth of a broad range of human tumor cell types, including clinical targets. Furthermore, recombinant adenoviruses can express tumor suppression genes such as p53 in an in vivo established tumor without relying on direct injection into the tumor or prior ex vivo treatment of the cancer cells. The p53 expressed is

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functional and effectively suppresses tumor growth *in vivo* and significantly increases survival time in a nude mouse model of human lung cancer.

the vectors of this Thus. invention are 5 particularly suited for gene therapy. Accordingly, methods of gene therapy utilizing these vectors are within the scope of this invention. The vector is purified and then an effective amount is administered in vivo or ex vivo into the subject. Methods of gene therapy are well known in the 10 art, see, for example, Larrick, J.W. and Burck, K.L. (1991) "Subject" means any animal, and Kreigler, M. (1990). mammal, rat, murine, bovine, porcine, equine, canine, feline or human patient. When the foreign gene codes for a tumor suppressor gene or other anti-tumor protein, the 15 vector is useful to treat or reduce hyperproliferative cells in a subject, to inhibit tumor proliferation in a subject or to ameliorate a particular related pathology. Pathologic hyperproliferative cells are characteristic of the following disease states, thyroid hyperplasia - Grave's 20 Disease, psoriasis, benign prostatic hypertrophy, Li-Fraumeni syndrome including breast cancer, sarcomas and other neoplasms, bladder cancer, colon cancer, lung cancer, various leukemias and lymphomas. Examples of nonhyperproliferative cells pathologic are found, for instance, in mammary ductal epithelial cells during development of lactation and also in cells associated with repair. Pathologic hyperproliferative cells characteristically exhibit loss of contact inhibition and a decline in their ability to selectively adhere which implies a change in the surface properties of the cell and a further breakdown in intercellular communication. These changes include stimulation to divide and the ability to secrete proteolytic enzymes.

Moreover, the present invention relates to a 35 method for depleting a suitable sample of pathologic

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hyperproliferative cells mammalian contaminating hematopoietic precursors during bone marrow reconstitution via the introduction of a wild type tumor suppressor gene into the cell preparation using the vector of this 5 invention (whether derived from autologous peripheral blood or bone marrow). As used herein, a "suitable sample" is defined as a heterogeneous cell preparation obtained from a patient, e.g., a mixed population of cells containing phenotypically normal and pathogenic 10 "Administer" includes, but is not limited to introducing into the cell or subject intravenously, by direct injection intra-tumoral tumor, рA injection, intraperitoneal administration, by aerosol administration to the lung or topically. Such administration can be 15 combined with a pharmaceutically-accepted carrier. described above.

The term "reduced tumorigenicity" is intended to mean tumor cells that have been converted into less tumorigenic or non-tumorigenic cells. Cells with reduced tumorigenicity either form no tumors in vivo or have an extended lag time of weeks to months before the appearance of in vivo tumor growth and/or slower growing three dimensional tumor mass compared to tumors having fully inactivated or non-functional tumor suppressor gene.

As used herein, the term "effective amount" is intended to mean the amount of vector or anti-cancer protein which achieves a positive outcome on controlling cell proliferation. For example, one dose contains from about 10⁸ to about 10¹³ infectious units. A typical course of treatment would be one such dose a day over a period of five days. An effective amount will vary on the pathology or condition to be treated, by the patient and his status, and other factors well known to those of skill in the art. Effective amounts are easily determined by those of skill in the art.

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Also within the scope of this invention is a method of ameliorating a pathology characterized by hyperproliferative cells or genetic defect in a subject by administering to the subject an effective amount of a vector described above containing a foreign gene encoding a gene product having the ability to ameliorate the pathology, under suitable conditions. As used herein, the term "genetic defect" means any disease or abnormality that results from inherited factors, such as sickle cell anemia or Tay-Sachs disease.

This invention also provides a method for reducing the proliferation of tumor cells in a subject by introducing into the tumor mass an effective amount of an adenoviral expression vector containing an anti-tumor gene 15 other than a tumor suppressor gene. The anti-tumor gene can encode, for example, thymidine kinase (TK). subject is then administered an effective amount of a therapeutic agent, which in the presence of the anti-tumor gene is toxic to the cell. In the specific case of 20 thymidine kinase, the therapeutic agent is a thymidine metabolite such as ganciclovir (GCV), methoxypurine arabinonucleoside (araM), or a functional equivalent thereof. Both the thymidine kinase gene and the thymidine kinase metabolite must be used concurrently to be 25 toxic to the host cell. However, in its presence, GCV is phosphorylated and becomes a potent inhibitor of DNA synthesis whereas araM gets converted to the cytotoxic anabolite araATP. Other anti-tumor genes can be used as well in combination with the corresponding therapeutic 30 agent to reduce the proliferation of tumor cells. other gene and therapeutic agent combinations are known by one skilled in the art. Another example would be the vector of this invention expressing the enzyme cytosine deaminase. Such vector would be used in conjunction with 35 administration of the drug 5-fluorouracil (Austin and Huber, 1993), or the recently described E. Coli Deo Δ gene

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in combination with 6-methyl-purine-2'-deosribonucleoside (Sorscher et al 1994).

As with the use of the tumor suppressor genes described previously, the use of other anti-tumor genes, 5 either alone or in combination with the appropriate therapeutic agent provides a treatment for the uncontrolled cell growth or proliferation characteristic of tumors and malignancies. Thus, this invention provides a therapy to stop the uncontrolled cellular growth in the patient 10 thereby alleviating the symptoms of the disease or cachexia present in the patient. The effect of this treatment includes, but is not limited to, prolonged survival time of the patient, reduction in tumor mass or burden, apoptosis of tumor cells or the reduction of the number of 15 circulating tumor cells. Means of quantifying the beneficial effects of this therapy are well known to those of skill in the art.

The invention provides a recombinant adenovirus expression vector characterized by the partial or total deletion of the adenoviral protein IX DNA and having a foreign gene encoding a foreign protein, wherein the foreign protein is a suicide gene or functional equivalent thereof. The anti-cancer gene TK, described above, is an example of a suicide gene because when expressed, the gene product is, or can be made to be lethal to the cell. For TK, lethality is induced in the presence of GCV. The TK gene is derived from herpes simplex virus by methods well known to those of skill in the art. The plasmid pMLBKTK in E. coli HB101 (from ATCC #39369) is a source of the herpes simplex virus (HSV-1) thymidine kinase (TK) gene for use in this invention. However, many other sources exist as well.

The TK gene can be introduced into the tumor mass by combining the adenoviral expression vector with a

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suitable pharmaceutically acceptable carrier. Introduction can be accomplished by, for example, direct injection of the recombinant adenovirus into the tumor mass. specific case of a cancer such as hepatocellular carcinoma 5 (HCC), direct injection into the hepatic artery can be used for delivery because most HCCs derive their circulation from this artery. To control proliferation of the tumor, cell death is induced by treating the patients with a TK metabolite such as ganciclovir to achieve reduction of The TK metabolite can be administered, for 10 tumor mass. example, systemically, by local innoculation into the tumor or in the specific case of HCC, by injection into the hepatic artery. The ΤK metabolite is preferably administered at least once daily but can be increased or 15 decreased according to the need. The TK metabolite can be administered simultaneous or subsequent to the administration of the TK containing vector. Those skilled in the art know or can determine the dose and duration which is therapeutically effective.

A method of tumor-specific delivery of a tumor suppressor gene is accomplished by contacting target tissue in an animal with an effective amount of the recombinant adenoviral expression vector of this invention. The gene is intended to code for an anti-tumor agent, such as a functional tumor suppressor gene or suicide gene. "Contacting" is intended to encompass any delivery method for the efficient transfer of the vector, such as intratumoral injection.

The use of the adenoviral vector of this invention to prepare medicaments for the treatment of a disease or for therapy is further provided by this invention.

The following examples are intended to illustrate, not limit the scope of this invention.

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EXPERIMENT NO. I

Plasmid pAd/MLP/p53/E1b- was used as the starting material for these manipulations. This plasmid is based on the pBR322 derivative pML2 (pBR322 deleted for base pairs 5 1140 to 2490) and contains adenovirus type 5 sequences extending from base pair 1 to base pair 5788 except that it is deleted for adenovirus type 5 base pairs 357 to 3327. At the site of the Ad5 357/3327 deletion a transcriptional unit is inserted which is comprised of the adenovirus type 10 2 major late promoter, the adenovirus type 2 tripartite leader cDNA and the human p53 cDNA. It is a typical E1 replacement vector deleted for the Ad5 E1a and E1b genes but containing the Ad5 protein IX gene (for review of Adenovirus vectors see: Graham and Prevec (1992)). 15 DNA was obtained from Gibco BRL. Restriction endonucleases and T4 DNA ligase were obtained from New England Biolabs. E. coli DH5lpha competent cells were purchased from Gibco BRL and 293 cells were obtained from the American Type Culture Collection (ATCC). Prep-A-Gene DNA purification resin was 20 obtained from BioRad. LB broth bacterial growth medium was obtained from Difco. Qiagen DNA purification columns were obtained from Qiagen, Inc. Ad5 dl327 was obtained from R.J. Schneider, NYU. The MBS DNA transfection kit was purchased from Stratagene.

25 One (1) μ g pAd/MLP/p53/E1b- was digested with 20 units each of restriction enzymes Ecl 136II and NgoMI according to the manufacturer's recommendations. Five (5) μ g Ad2 DNA was digested with 20 units each of restriction endonucleases DraI and NgoMI according the 30 manufacturer's recommendations. The restriction digestions were loaded into separate lanes of a 0.8% agarose gel and electrophoresed at 100 volts for 2 hours. The 4268 bp restriction fragment from the Pad/MLP/p53/E1b- sample and the 6437 bp fragment from the Ad2 sample were isolated from 35 the gel using Prep-A-Gene DNA extraction resin according to

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manufacturer's specifications. The restriction fragments were mixed and treated with T4 DNA ligase in a total volume of 50 μ l at 16°C for 16 hours according to the manufacturer's recommendations. Following ligation 5 μ l of 5 the reaction was used to transform $E.\ coli$ DH5 α cells to ampicillin resistance following the manufacturer's Six bacterial colonies resulting from this procedure. procedure were used to inoculate separate 2 ml cultures of LB growth medium and incubated overnight at 37°C with 10 shaking. DNA was prepared from each bacterial culture using standard procedures (Sambrook et al. (1989)). fourth of the plasmid DNA from each isolate was digested with 20 units of restriction endonuclease XhoI to screen for the correct recombinant containing XhoI restriction 15 fragments of 3627, 3167, 2466 and 1445 base pairs. Five of six screened isolates contained the correct plasmid. of these was then used to inoculate a 1 liter culture of LB medium for isolation of large quantities of plasmid DNA. Following overnight incubation plasmid DNA was isolated 20 from the 1 liter culture using Qiagen DNA purification columns according to the manufacturer's recommendations. The resulting plasmid was designated Pad/MLP/p53/PIX-. Samples of this plasmid were deposited with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland, U.S.A., 12301, on October 22, 1993. The deposit was made under the provisions of the Budapest Treaty on the International Deposit of Microorganisms for the Purpose of Patent Procedure. The deposit was accorded ATCC Accession No. 75576.

To construct a recombinant adenovirus, 10 μ g Pad/MLP/p53/PIX- were treated with 40 units of restriction endonuclease EcoRI to linearize the plasmid. Adenovirus type 5 dl327 DNA (Thimmappaya (1982)) was digested with restriction endonuclease ClaI and the large fragment (approximately 33 kilobase pairs) was purified by sucrose gradient centrifugation. Ten (10) μ g of EcoRI treated

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Pad/MLP/p53/E1b- and 2.5 μg of ClaI treated Ad5 dl327 were mixed and used to transfect approximately 10⁶ 293 cells using the MBS mammalian transfection kit as recommended by the supplier. Eight (8) days following the transfection the 293 cells were split 1 to 3 into fresh media and two days following this adenovirus induced cytopathic effect became evident on the transfected cells. At 13 days post-transfection DNA was prepared from the infected cells using standard procedures (Graham and Prevec (1991)) and analyzed by restriction digestion with restriction endonuclease XhoI. Virus directed expression of p53 was verified following infection of SaoS2 osteosarcoma cells with viral lysate and immunoblotting with an anti-p53 monoclonal antibody designated 1801 (Novocasta Lab. Ltd., U.K.).

15 EXPERIMENT NO. II

MATERIALS AND METHODS

Cell Lines

Recombinant adenoviruses were grown and propagated in the human embryonal kidney cell line 293
20 (ATCC CRL 1573) maintained in DME medium containing 10% defined, supplemented calf serum (Hyclone). Saos-2 cells were maintained in Kaighn's media supplemented with 15% fetal calf serum. HeLa and Hep 3B cells were maintained in DME medium supplemented with 10% fetal calf serum. All other cell lines were grown in Kaighn's media supplemented with 10% fetal calf serum. Saos-2 cells were kindly provided by Dr. Eric Stanbridge. All other cell lines were obtained from ATCC.

Construction of Recombinant Adenoviruses

30 To construct the Ad5/p53 viruses, a 1.4 kb HindIII-SmaI fragment containing the full length cDNA for

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p53 (Table I) was isolated from pGEM1-p53-B-T (kindly supplied by Dr. Wen Hwa Lee) and inserted into the multiple cloning site of the expression vector pSP72 (Promega) using standard cloning procedures (Sambrook et al. (1989)). 5 p53 insert was recovered from this vector following digestion with XhoI-BglII and gel electrophoresis. The p53 coding sequence was then inserted into either pNL3C or pNL3CMV adenovirus gene transfer vectors (kindly provided by Dr. Robert Schneider) which contain the Ad5 5' inverted 10 terminal repeat and viral packaging signals and the Ela enhancer upstream of either the Ad2 major late promoter (MLP) or the human cytomegalovirus immediate early gene promoter (CMV), followed by the tripartite leader CDNA and Ad 5 sequence 3325-5525 bp in a PML2 background. These new 15 constructs replace the E1 region (bp 360-3325) of Ad5 with p53 driven by either the Ad2 MLP (A/M/53) or the human CMV promoter (A/C/53), both followed by the tripartite leader CDNA (see Figure 4). The p53 inserts use the remaining downstream Elb polyadenylation site. Additional MLP and 20 CMV driven p53 recombinants (A/M/N/53, A/C/N/53) were generated which had a further 705 nucleotide deletion of Ad 5 sequence to remove the protein IX (PIX) coding region. As a control, a recombinant adenovirus was generated from the parental PNL3C plasmid without a p53 insert (A/M). 25 second control consisted of a recombinant adenovirus encoding the beta-galactosidase gene under the control of the CMV promoter (A/C/ß-gal). The plasmids were linearized with either Nru I or Eco RI and co-transfected with the large fragment of a Cla I digested Ad 5 d1309 or d1327 30 mutants (Jones and Shenk (1979)) using a Ca/PO4 transfection (Stratagene). Viral plaques were isolated and recombinants identified by both restriction digest analysis and PCR using recombinant specific primers against the tripartite leader CDNA sequence with downstream p53 CDNA Recombinant virus was further purified by 35 sequence. limiting dilution, and virus particles were purified and

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titered by standard methods (Graham and van der Erb (1973); Graham and Prevec (1991)).

p53 Protein Detection

Saos-2 or Hep 3B cells (5 \times 10 5) were infected 5 with the indicated recombinant adenoviruses for a period of 24 hours at increasing multiplicities of infection (MOI) of plaque forming units of virus/cell. Cells were then washed once with PBS and harvested in lysis buffer (50mM Tris-Hcl Ph 7.5, 250 Mm NaCl, 0.1% NP40, 50mM NaF, 5mM EDTA, 10ug/ml 10 aprotinin, 10 ug/ml leupeptin, and 1mM PMSF). proteins (approximately 30 μg) were separated by 10% SDS-PAGE and transferred to nitrocellulose. Membranes were incubated with α -p53 antibody PAb 1801 (Novocastro) followed by sheep anti-mouse IgG conjugated with 15 horseradish peroxidase. p53 protein was visualized by chemiluminescence (ECL kit, Amersham) on Kodak XAR-5 film.

Measurement of DNA Synthesis Rate

Cells (5 x 10³/well) were plated in 96-well titer plates (Costar) and allowed to attach overnight (37°C, 7% CO₂). Cells were then infected for 24 hours with purified recombinant virus particles at MOIs ranging from 0.3 to 100 as indicated. Media were changed 24 hours after infection, and incubation was continued for a total of 72 hours. 3 H-thymidine (Amersham, 1 μ Ci/well) was added 18 hours prior to harvest. Cells were harvested on glass fiber filters and levels of incorporated radioactivity were measured in a beta scintillation counter. 3 H-thymidine incorporation was expressed as the mean % (+/- SD) of media control and plotted versus the MOI.

30 Tumorigenicity in Nude Mice

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Approximately 2.4 x 108 Saos-2 cells, plated in T225 flasks, were treated with suspension buffer (1% sucrose in PBS) containing either A/M/N/53 or A/M purified virus at an MOI of 3 or 30. Following an overnight infection, cells were injected subcutaneously into the left and right flanks of BALB/c athymic nude mice (4 mice per group). One flank was injected with the A/M/N/53 treated cells, while the contralateral flank was injected with the control A/M treated cells, each mouse serving as its own 10 control. Animals receiving bilateral injection of buffer treated cells served as additional controls. dimensions (length, width and height) and body weights were then measured twice per week over an 8 week period. Tumor volumes were estimated for each animal assuming a spherical 15 geometry with radius equal to one-half the average of the measured tumor dimensions.

Intra-tumoral RNA Analysis

BALB/c athymic nude mice (approximately 5 weeks of age) were injected subcutaneously with 1 x 10^7 H69 small cell lung carcinoma (SCLC) cells in their right flanks. 20 Tumors were allowed to progress for 32 days until they were approximately 25-50 mm³. Mice received peritumoral injections of either A/C/53 or A/C/B-gal recombinant adenovirus (2 x 109 plaque forming units (pfu)) into the 25 subcutaneous space beneath the tumor mass. Tumors were excised from the animals 2 and 7 days post adenovirus treatment and rinsed with PBS. Tumor samples were homogenized, and total RNA was isolated using a TriReagent kit (Molecular Research Center, Inc.). PolyA RNA was isolated using the PolyATract mRNA Isolation System 30 (Promega), and approximately 10 ng of sample was used for RT-PCR determination of recombinant p53 MRNA expression (Wang et al. (1989)). Primers were designed to amplify sequence between the adenovirus tripartite leader CDNA and

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the downstream p53 CDNA, ensuring that only recombinant, and not endogenous p53 would be amplified.

p53 Gene Therapy of Established Tumors in Nude Mice

Approximately 1 x 10⁷ H69 (SCLC) tumor cells in 200µl volumes were injected subcutaneously into female BALB/c athymic nude mice. Tumors were allowed to develop for 2 weeks, at which point animals were randomized by tumor size (N=5/group). Peritumoral injections of either A/M/N/53 or the control A/M adenovirus (2 x 10⁹ pfu/injection) or buffer alone (1% sucrose in PBS) were administered twice per week for a total of 8 doses/group. Tumor dimensions and body weights were measured twice per week for 7 weeks, and tumor volume was estimated as described previously. Animals were then followed to observe the effect of treatment on mouse survival.

RESULTS

Construction of Recombinant p53-Adenovirus

p53 adenoviruses were constructed by replacing a portion of the Ela and Elb region of adenovirus Type 5 with p53 CDNA under the control of either the Ad2 MLP (A/M/53) or CMV (A/C/53) promoter (schematized in Figure 4). This El substitution severely impairs the ability of the recombinant adenoviruses to replicate, restricting their propagation to 293 cells which supply Ad 5 El gene products in trans (Graham et al. (1977)). After identification of p53 recombinant adenovirus by both restriction digest and PCR analysis, the entire p53 CDNA sequence from one of the recombinant adenoviruses (A/M/53) was sequenced to verify that it was free of mutations. Following this, purified preparations of the p53 recombinants were used to infect HeLa cells to assay for the presence of phenotypically wild type adenovirus. HeLa cells, which are non-permissive for

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replication of E1-deleted adenovirus, were infected with 1-4 x 10° infectious units of recombinant adenovirus, cultured for 3 weeks, and observed for the appearance of cytopathic effect (CPE). Using this assay, recombinant adenovirus replication or wild type contamination was not detected, readily evident by the CPE observed in control cells infected with wild type adenovirus at a level of sensitivity of approximately 1 in 10°.

p53 Protein Expression from Recombinant Adenovirus

10 To determine if p53 recombinant adenoviruses expressed p53 protein, tumor cell lines which do not express endogenous p53 protein were infected. The human cell lines Saos-2 (osteosarcoma) and Hep (hepatocellular carcinoma) were infected for 24 hours with 15 the p53 recombinant adenoviruses A/M/53 or A/C/53 at MOIs ranging 0.1 to 200 pfu/cell. Western analysis of lysates prepared from infected cells demonstrated a dose-dependent p53 protein expression in both cell types (Figure 5). Both cell lines expressed higher levels of p53 protein following 20 infection with A/C/53 than with A/M/53 (Figure 3). No p53 protein was detected in non-infected cells. endogenous wild-type p53 are normally quite low, and nearly undetectable by Western analysis of cell extracts (Bartek et al. (1991)). It is clear however that wild-type p53 25 protein levels are easily detectable after infection with either A/M/53 or A/C/53 at the lower MOIs (Figure 5), suggesting that even low doses of p53 recombinant adenoviruses can produce potentially efficacious levels of p53.

30 p53 Dependent Morphology Changes

The reintroduction of wild-type p53 into the p53-negative osteosarcoma cell line, Saos-2, results in a characteristic enlargement and flattening of these normally

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spindle-shaped cells (Chen et al. (1990)). Subconfluent Saos-2 cells (1x105 cells/10cm plate) were infected at an MOI of 50 with either the A/C/53 or control A/M virus, and incubated at 37°C for 72 hours until uninfected control 5 plates were confluent. At this point, the expected morphological change was evident in the A/C/53 treated plate (Figure 6, panel C) but not in uninfected (Figure 6, panel A) or control virus-infected plates (Figure 6, panel B). This effect was not a function of cell density because 10 a control plate initially seeded at lower density retained normal morphology at 72 hours when its confluence approximated that of the A/C/53 treated plate. results had demonstrated a high level of p53 protein expression at an MOI of 50 in Saos-2 cells (Figure 5A), and 15 these results provided evidence that the p53 protein expressed by these recombinant adenoviruses was biologically active.

p53 Inhibition of Cellular DNA Synthesis

To further test the activity of the 20 recombinant adenoviruses, their ability to proliferation of human tumor cells was assayed as measured by the uptake of ³H-thymidine. It has previously been shown that introduction of wild-type p53 into cells which do not express endogenous wild-type p53 can arrest the cells at the G₁/S transition, leading to inhibition of uptake of labeled thymidine into newly synthesized DNA (Baker et al. (1990); Mercer et al. (1990); Diller et al. (1990)). variety of p53-deficient tumor cell lines were infected with either A/M/N/53, A/C/N/53 or a non-p53 expressing 30 control recombinant adenovirus (A/M). A strong, dosedependent inhibition of DNA synthesis by both the A/M/N/53and A/C/N/53 recombinants in 7 out of the 9 different tumor cell lines tested (Figure 7) was observed. Both constructs were able to inhibit DNA synthesis in these human tumor 35 cells, regardless of whether they expressed mutant p53 or

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failed to express p53 protein. It also was found that in this assay, the A/C/N/53 construct was consistently more potent than the A/M/N/53. In saos-2 (osteosarcoma) and MDA-MB468 (breast cancer) cells, nearly 100% inhibition of DNA synthesis was achieved with the A/C/N/53 construct at an MOI as low as 10. At doses where inhibition by the control adenovirus in only 10-30%, a 50-100% reduction in DNA synthesis using either p53 recombinant adenovirus was observed. In contrast, no significant p53-specific effect was observed with either construct as compared to control virus in HEP G2 cells (hepatocarcinoma cell line expressing endogenous wild-type p53, Bressac et al. (1990)), nor in the K562 (p53 null) leukemic cell line.

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Tumorigenicity in Nude Mice

In a more stringent test of function for the p53 recombinant adenoviruses, tumor cells were infected ex vivo and then injected the cells into nude mice to assess the 5 ability of the recombinants to suppress tumor growth in vivo. Saos-2 cells infected with A/M/N/53 or control A/M virus at a MOI of 3 or 30, were injected into opposite flanks of nude mice. Tumor sizes were then measured twice a week over an 8 week period. At the MOI of 30, no tumor 10 growth was observed in the p53-treated flanks in any of the animals, while the control treated tumors continued to grow (Figure 8). The progressive enlargement of the control virus treated tumors were similar to that observed in the buffer treated control animals. A clear difference in 15 tumor growth between the control adenovirus and the p53 recombinant at the MOI of 3, although tumors from 2 out of the 4 p53-treated mice did start to show some growth after approximately 6 weeks. Thus, the A/M/N/53 recombinant adenovirus is able to mediate p53-specific 20 suppression in an in vivo environment.

In Vivo Expression of Ad/p53

Although ex vivo treatment of cancer cells and subsequent injection into animals provided a critical test of tumor suppression, a more clinically relevant experiment is to determine if injected p53 recombinant adenovirus could infect and express p53 in established tumors in vivo. To address this, H69 (SCLC, p53^{null}) cells were injected subcutaneously into nude mice, and tumors were allowed to develop for 32 days. At this time, a single injection of 2 x 10⁹ pfu of either A/C/53 or A/C/ß-gal adenovirus was injected into the peritumoral space surrounding the tumor. Tumors were then excised at either Day 2 or Day 7 following the adenovirus injection, and polyA RNA was isolated from each tumor. RT-PCR, using recombinant-p53 specific

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primers, was then used to detect p53 MRNA in the p53 treated tumors (Figure 9, lanes 1,2,4,5). No p53 signal was evident from the tumors excised from the ß-gal treated animals (Figure 9, lanes 3 and 6). Amplification with actin primers served as a control for the RT-PCR reaction (Figure 9, lanes 7-9), while a plasmid containing the recombinant-p53 sequence served as a positive control for the recombinant-p53 specific band (Figure 9, lane 10). This experiment demonstrates that a p53 recombinant adenovirus can specifically direct expression of p53 mRNA within established tumors following a single injection into the peritumoral space. It also shows in vivo viral persistence for at least one week following infection with a p53 recombinant adenovirus.

15 In Vivo Efficacy

To address the feasibility of gene therapy of established tumors, a tumor-bearing nude mouse model was used. H69 cells were injected into the subcutaneous space on the right flank of mice, and tumors were allowed to grow 20 for 2 weeks. Mice then received peritumoral injections of buffer or recombinant virus twice weekly for a total of 8 In the mice treated with buffer or control A/M doses. virus, tumors continued to grow rapidly throughout the treatment, whereas those treated with the A/M/N/53 virus 25 grew at a greatly reduced rate (Figure 10A). cessation of injections, the control treated tumors continued to grow while the p53 treated tumors showed little or no growth for at least one week in the absence of any additional supply of exogenous p53 (Figure 10A). Although control animals treated with buffer alone had 30 accelerated tumor growth as compared to either virus treated group, no significant difference in body weight was found between the three groups during the treatment period. Tumor ulceration in some animals limited the relevance of 35 tumor size measurements after day 42. However, continued

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monitoring of the animals to determine survival time demonstrated a survival advantage for the p53-treated animals (Figure 10B). The last of the control adenovirus treated animals died on day 83, while buffer alone treated controls had all expired by day 56. In contrast, all 5 animals treated with the A/M/N/53 continue to survive (day 130 after cell inoculation) (Figure 10B). Together, this data establish a p53-specific effect on both tumor growth and survival time in animals with established p53-deficient tumors.

Adenovirus Vectors Expressing p53

Recombinant human adenovirus vectors which are capable of expressing high levels of wild-type p53 protein in a dose dependent manner were constructed. 15 contains deletions in the Ela and Elb regions which render the virus replication deficient (Challberg and Kelly (1979); Horowitz, (1991)). Of further significance is that these deletions include those sequences encoding the E1b 19 and 55 kd protein. The 19 kd protein is reported to be 20 involved in inhibiting apoptosis (White et al. (1992); Rao et al. (1992)), whereas the 55 kd protein is able to bind wild-type p53 protein (Sarnow et al. (1982); Heuvel et al. (1990)). By deleting these adenoviral sequences, potential inhibitors of p53 function were removed through direct 25 binding to p53 or potential inhibition of p53 mediated apoptosis. Additional constructs were made which have had the remaining 3' Elb sequence, including all protein IX coding sequence, deleted as well. Although this has been reported to reduce the packaging size capacity of adenovirus to approximately 3 kb less than wild-type virus 30 (Ghosh-Choudhury et al. (1987)), these constructs are also deleted in the E3 region so that the A/M/N/53 and A/C/N/53constructs are well within this size range. By deleting the pIX region, adenoviral sequences homologous to those 35 contained in 293 cells are reduced to approximately 300

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base pairs, decreasing the chances of regenerating replication-competent, wild-type adenovirus through recombination. Constructs lacking pIX coding sequence appear to have equal efficacy to those with pIX.

5 p53/Adenovirus Efficacy In Vitro

In concordance with a strong dose dependency for expression of p53 protein in infected cells, a dosedependent, p53-specific inhibition of tumor cell growth was demonstrated. Cell division. was inhibited 10 demonstrated by the inhibition of DNA synthesis, in a wide variety of tumor cell types known to lack wild-type p53 protein expression. Bacchetti and Graham (1993) recently reported p53 specific inhibition of DNA synthesis in the ovarian carcinoma cell line SKOV-3 by a p53 recombinant 15 adenovirus in similar experiments. In addition to ovarian carcinoma, additional human tumor cell lines demonstrated, representative of clinically important human cancers and including lines over-expressing mutant p53 protein, can also be growth inhibited by the 20 recombinants of this invention. At MOIs where the A/C/N/53recombinant is 90-100% effective in inhibiting synthesis in these tumor types, control adenovirus mediated suppression is less than 20%.

Although Feinstein et al. (1992) reported that 25 re-introduction of wild-type p53 could differentiation and increase the proportion of cells in G_1 versus S+G2 for leukemic K562 cells, no p53 specific effect was found in this line. Horvath and Weber (1988) have reported that human peripheral blood lymphocytes are highly nonpermissive to adenovirus infection. In separate experiments, the recombinant significantly infected the non-responding K562 cells with recombinant A/C/ß-gal adenovirus, while other cell lines, including the control Hep G2 line and those showing a strong p53 effect, were

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readily infectable. Thus, at least part of the variability of efficacy would appear to be due to variability of infection, although other factors may be involved as well.

The results observed with the A/M/N/53 virus in 5 Figure 8 demonstrates that complete suppression is possible in an *in vivo* environment. The resumption of tumor growth in 2 out of 4, p53 treated animals at the lower MOI most likely resulted from a small percentage of cells not initially infected with the p53 recombinant at this dose.

10 The complete suppression seen with A/M/N/53 at the higher dose, however, shows that the ability of tumor growth to recover can be overcome.

p53/Adenovirus In Vivo Efficacy

Work presented here and by other groups (Chen et 15 al. (1990); Takahashi et al. (1992)) have shown that human tumor cells lacking expression of wild-type p53 can be treated ex vivo with p53 and result in suppression of tumor growth when the treated cells are transferred into an animal model. Applicants present the first evidence of 20 tumor suppressor gene therapy of an in vivo established tumor, resulting in both suppression of tumor growth and increased survival time. In Applicants' system, delivery to tumor cells did not rely on direct injection into the tumor mass. Rather, p53 recombinant adenovirus was 25 injected into the peritumoral space, and p53 mRNA expression was detected within the tumor. p53 expressed by the recombinants was functional and strongly suppressed tumor growth as compared to that of control, non-p53 expressing adenovirus treated tumors. However, both p53 30 and control virus treated tumor groups showed tumor suppression as compared to buffer treated controls. It has been demonstrated that local expression of tumor necrosis factor (TNF), interferon- γ), interleukin (IL)-2, IL-4 or IL-7 can lead to T-cell independent transient tumor

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suppression in nude mice (Hoch et al. (1992)). Exposure of monocytes to adenovirus virions are also weak inducers of IFN- α/β (reviewed in Gooding and Wold (1990)). Therefore, it is not surprising that some tumor suppression in nude 5 mice was observed even with the control adenovirus. virus mediated tumor suppression was not observed in the ex vivo control virus treated Saos-2 tumor cells described The p53-specific in vivo tumor suppression was dramatically demonstrated by continued monitoring of the 10 animals in Figure 10. The survival time of the p53-treated mice was significantly increased, with 5 out of 5 animals still alive more than 130 days after cell inoculation compared to 0 out of 5 adenovirus control treated animals. The surviving animals still exhibit growing tumors which 15 may reflect cells not initially infected with the p53 recombinant adenovirus. Higher or more frequent dosing schedules may address this. In addition, promoter shutoff (Palmer et al. (1991)) or additional mutations may have rendered these cells resistant to the p53 recombinant 20 adenovirus treatment. For example, mutations in the recently described WAF1 gene, a gene induced by wild-type p53 which subsequently inhibits progression of the cell cycle into S phase, (El-Deiry et al. (1993); Hunter (1993)) could result in a p53-resistant tumor.

25 EXPERIMENT NO. III

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This Example shows the use of suicide genes and tissue specific expression of such genes in the gene therapy methods described herein. Hepatocellular carcinoma was chosen as the target because it is one of the most common human malignancies affecting man, causing estimated 1,250,000 deaths per year world-wide. incidence of this cancer is very high in Southeast Asia and Africa where it is associated with Hepatitis B and C infection and exposure to aflatoxin. Surgery is currently 35 the only treatment which offers the potential for curing

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HCC, although less than 20% of patients are considered candidates for resection (Ravoet C. et al., 1993). However, tumors other than hepatocellular carcinoma are equally applicable to the methods of reducing their proliferation described herein.

CELL LINES

All cell lines but for the HLF cell line were obtained from the American Type Tissue Culture Collection (ATCC) 12301 Parklawn Drive, Rockville Maryland. accession numbers are noted in parenthesis. The human embryonal kidney cell line 293 (CRL 1573) was used to generate and propagate the recombinant adenoviruses They were maintained in DME medium described herein. containing 10% defined, supplemented calf serum (Hyclone). The hepatocellular carcinoma cell lines Hep 3B (HB 8064), Hep G2 (HB 8065), and HLF were maintained in DME/F12 medium supplemented with 10% fetal bovine serum, as were the breast carcinoma cell lines MDA-MB468 (HTB 132) and BT-549 (HTB 122). Chang liver cells (CCL 13) were grown in MEM 20 medium supplemented with 10% fetal bovine serum. The HLF cell line was obtained from Drs. T. Morsaki and H. Kitsuki at the Kyushu University School of Medicine in Japan.

RECOMBINANT VIRUS CONSTRUCTION

Two adenoviral expression vectors designated 25 herein as ACNTK and AANTK and devoid of protein IX function (depicted in Figure 11) are capable of directing expression of the TK suicide gene within tumor cells. adenovirus expression vector designated AANCAT constructed to further demonstrate the feasibility of 30 specifically targeting gene expression to specific cell types using adenoviral vectors. These adenoviral constructs were assembled as depicted in Figures 11 and 12 and are derivatives of those previously described for the expression of tumor suppresor genes.

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For expression of the foreign gene, expression cassettes have been inserted that utilize either the human cytomegalovirus immediate early promoter/enhancer (CMV) (Boshart, M. et al., 1985) or the human alpha-fetoprotein 5 (AFP) enhancer/promoter (Watanable, K. et al., Nakabayashi, H. et al., 1989) to direct transcription of the TK gene or the chloramphenicol acetyltransferase gene The CMV enhancer promoter is capable of directing robust gene expression in a wide variety of cell types 10 while the enhancer/promoter construct restricts AFP expression to hepatocellular carcinoma cells (HCC) which express AFP in about 70-80% of the HCC pateint population. In the construct utilizing the CMV promoter/enhancer, the adenovirus type 2 tripartite leader sequence also was 15 inserted to enhance translation of the TK transcript (Berkner, K.L. and Sharp, 1985). In addition to the E1 deletion, both adenovirus vectors are additionally deleted for 1.9 kilobases (kb) of DNA in the viral E3 region. DNA deleted in the E3 region is non-essential for virus 20 propagation and its deletion increases the insert capacity of the recombinant virus for foreign DNA by an equivalent amount (1.9kb) (Graham and Prevec, 1991).

demonstrate the specificity of the AFP promoter/enhancer, the virus AANCAT also was constructed 25 where the marker gene chloramphenicol aceytitransferase (CAT) is under the control of the AFP enhancer/promoter. In the ACNTK viral construct, the Ad2 tripartite leader sequence was placed between the CMV promoter/enhancer and the TK gene. The tripartite leader has been reported to enhance translation of linked genes. The El substitution the ability of the recombinant viruses replicate, restricting their propagation to 293 cells which supply the Ad5 E1 gene products in trans (Graham et al., 1977).

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Adenoviral Vector ACNTK: The plasmid pMLBKTK in E. coli HB101 (from ATCC #39369) was used as the source of the herpes simplex virus (HSV-1) thymidine kinase (TK) TK was excised from this plasmid as a 1.7 kb gene 5 fragment by digestion with the restriction enzymes Bgl II and Pvu II and subcloned into the compatible Bam HI, EcoR V restriction sites of plasmid pSP72 (Promega) using standard cloning techniques (Sambrook et al., 1989). TK insert was then isolated as a 1.7 kb fragment from this 10 vector by digestion with Xba I and Bgl II and cloned into Xba I, BamHI digested plasmid pACN (Wills et al. 1994). Twenty (20) μ g of this plasmid designated pACNTK were linearized with Eco RI and cotransfected into 293 cells (ATCC CRL 1573) with 5 μg of Cla I digested ACBGL (Wills et al., 1994 supra) using a CaPO, transfection kit (Stratagene, San Diego, California). Viral plaques were isolated and recombinants, designated ACNTK, were identified restriction digest analysis of isolated DNA with Xho I and BsiWI. Positive recombinants were further purified by limiting dilution and expanded and titered by standard methods (Graham and Prevec, 1991).

Adenoviral Vector AANTK: The α -fetoprotein promoter (AFP-P) and enhancer (AFP-E) were cloned from a human genomic DNA (Clontech) using PCR amplification with 25 primers containing restriction sites at their ends. primers used to isolate the 210 bp AFP-E contained a Nhe I restriction site on the 5' primer and an Xba I, Xho I, Kpn I linker on the 3' primer. The 5' primer sequence was 5'-CGC GCT AGC TCT GCC CCA AAG AGC T-3. The 5' primer sequence was 5'-CGC GGT ACC CTC GAG TCT AGA TAT TGC CAG TGG The primers used to isolate the 1763 bp AFE TGG AAG-3'. fragment contained a Not I restriction site on the 5' primer and a Xba I site on the 3' primer. The 5' primer sequence was 5'-CGT GCG GCC GCT GGA GGA CTT TGA GGA TGT CTG TC-3'. The 3' primer sequence was 5'-CGC TCT AGA GAG ACC 35 AGT TAG GAA GTT TTC GCA-3'. For PCR amplification, the DNA

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was denatured at 97° for 7 minutes, followed by 5 cycles of amplification at 97°, 1 minute, 53°, 1 minute, 72°, 2 minutes, and a final 72°, 10 minute extension. The amplified AFE was digested with Not I and Xba I and inserted into the Not I, Xba I sites of a plasmid vector (pA/ITR/B) containing adenovirus type 5 sequences 1-350 and 3330 - 5790 separated by a polylinker containing Not I, Xho I, Xba I, Hind III, Kpn I, Bam HI, Nco I, Sma I, and Bgl II The amplified AFP-E was digested with Nhe I and Kpn and inserted into the AFP-E containing construct 10 described above which had been digested with Xba I and Kpn This new construct was then further digested with Xba I and NgoMI to remove adenoviral sequences 3330 - 5780, which were subsequently replaced with an Xba I, NgoMI 15 restriction fragment of plasmid pACN containing nucleotides 4021 - 10457 of adenovirus type 2 to construct the plasmid pAAN containing both the α-fetoprotein enhancer promoter. This construct was then digested with Eco RI and Xba I to isolate a 2.3 kb fragment containing the Ad5 20 inverted terminal repeat, the AFP-E and the AFP-P which was subsequently ligated with the 8.55 kb fragment of Eco RI, Xba I digested pACNTK described above to generate pAANTK where the TK gene is driven by the α -fetoprotein enhancer and promoter in an adenovirus background. This plasmid was 25 then linearized with Eco RI and cotransfected with the large fragment of Cla I digested ALBGL as above and recombinants, designated AANTK, were isolated and purified as described above.

Adenoviral Vector AANCAT: The chloramphenicol acetyltransferase (CAT) gene was isolated from the pCAT-Basic Vector (Promega Corporation) by an Xba I, Bam HI digest. This 1.64 kb fragment was ligated into Xba I, Bam HI digested pAAN (described above) to create pAANCAT. This plasmid was then linearized with Eco RI and cotransfected with the large fragment of Cla I digested rA/C/S-gal to create AANCAT.

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REPORTER GENE EXPRESSION: &-GALACTOSIDASE EXPRESSION:

Cells were plated at 1 x 10⁵ cells/well in a 24-well tissue culture plate (Costar) and allowed to adhere overnight (37C, 7% CO₂). Overnight infections of ACBGL were performed at a multiplicity of infection (MOI) of 30. After 24 hours, cells were fixed with 3.7% Formaldehyde; PBS, and stained with 1mg/ml Xgal reagent (USB). The data was scored (+,++,+++) by estimating the percentage of positively stained cells at each MOI. [+=1-33%, ++=33-67% and +++=>67%]

REPORTER GENE EXPRESSION: CAT EXPRESSION:

Two (2) x 106 cells (Hep G2, Hep 3B, HLF, Chang, and MDA-MB468) were seeded onto 10 cm plates in triplicate and incubated overnight (37C, 7% CO_2). Each plate was then infected with either AANCAT at an MOI = 30 or 100 or uninfected and allowed to incubate for 3 days. The cells were then trypsinized and washed with PBS and resuspended in 100 μ l of 0.25 M Tris pH 7.8. The samples were frozen and thawed 3 times, and the supernatant was transferred to new tubes and incubated at 60°C for 10 minutes. 20 samples were then spun at 4°C for 5 minutes, and the supernatants assayed for protein concentration using a Bradford assay (Bio-Rad Protein Assay Kit). Samples were adjusted to equal protein concentrations to a final volume 25 of 75 μ l using 0.25 M Tris, 25 μ l of 4mM acetyl CoA and 1 μ l of ¹⁴C-Chloramphenicol and incubated overnight at 37°C. 500 μ l of ethyl acetate is added to each sample and mixed by vortexing, followed by centrifiguration for 5 minutes at room temperature. The upper phase is then transferred to 30 a new tube and the ethyl acetate is evaporated by centrifugation under vacuum. The reaction products are then redissolved in 25 μ l of ethyl acetate and spotted onto a thin layer chromatography (TLC) plate and the plate is then placed in a pre-equilibrated TLC chamber (95%

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chloroform, 5% methanol). The solvent is then allowed to migrate to the top of the plate, the plate is then dried and exposed to X-ray film.

CELLULAR PROLIFERATION: 3H-THYMIDINE INCORPORATION

Cells were plated at 5 x 10³ cells/well in a 96-5 well micro-titer plate (Costar) and allowed to incubate overnight (37C, 7% CO₂). Serially diluted ACN, ACNTK or AATK virus in DMEM; 15% FBS; 1% glutamine was used to transfect cells at an infection multiplicity of 30 for an 10 overnight duration at which point cells were dosed in triplicate with ganciclovir (Cytovene) at log intervals betweem 0.001 and 100 mM (micro molar). 1 μ Ci 3 H-thymidine (Amersham) was added to each well 12-18 hours before harvesting. At 72 hours-post infection cells were 15 harvested onto glass-fiber filters and incorporated ³Hthymidine was counted using liquid scintillation (TopCount, Results are plotted as percent of untreated Packard). control proliferation and tabulated as the effective dose $(ED_{so}\pm SD)$ for a 50 percent reduction in proliferation over media controls. ED_{50} values were estimated by fitting a logistic equation to the dose response data.

CYTOTOXICITY: LDH RELEASE

Cells (HLF, human HCC) were plated, infected with ACN or ACNTK and treated with ganciclovir as described for the proliferation assay. At 72 hours post-ganciclovir administration, cells were spun, the supernatant was removed. The levels of lactate dehydrogenase measured colometrically (Promega, Cytotox 96TM). Mean (+/- S.D.) LDH release is plotted versus M.O.I.

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Human hepatocellular carcinoma cells (Hep 3B) were injected subcutaneously into ten female (10) athymic nu/nu mice (Simonsen Laboratories, Gilroy, CA). animal received approximately 1 x 107 cells in the left Tumors were allowed to grow for 27 days before 5 flank. randomizing mice by tumor size. Mice were treated with intratumoral and peritumoral injections of ACNTK or the control virus ACN (1 x 10^9 iu in 100μ l) every other day for a total of three doses. Starting 24 hours after the adenovirus, the 10 initial dose of mice were dosed intraperitoneally with ganciclovir (Cytovene 100 mg/kg) daily for a total of 10 days. Mice were monitored for tumor size and body weight twice weekly. Measurements on tumors were made in three dimensions using vernier calipers 15 and volumes were calculated using the formula $4/3 \pi r^3$, where r is one-half the average tumor dimension.

RESULTS

The recombinant adenoviruses were used to infect three HCC cell lines (HLF, Hep3B and Hep-G2). One human liver cell line (Chang) and two breast cancer cell lines were used as controls (MDAMB468 and BT549). To demonstrate the specificity of the AFP promoter/enhancer, the virus AANCAT was constructed. This virus was used to infect cells that either do (Hep 3B, HepG2) or do not (HLE, Chang, MDAMB468) express the HCC tumor marker alphafetoprotein (AFP). As shown in Figure 13, AANCAT directs expression of the CAT marker gene only in those HCC cells which are capable of expressing AFP (Figure 13).

The efficacy of ACNTK and AANTK for the treatment of HCC was assessed using a ³H-thymidine incorporation assay to measure the effect of the combination of HSV-TK expression and ganciclovir treatment upon cellular proliferation. The cell lines were infected with either ACNTK or AANTK or the control virus ACN (Wills et al., 1994

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supra), which does not direct expression of HSV-TK, and then treated with increasing concentrations of ganciclovir. The effect of this treatment was assessed as a function of increasing concentrations of ganciclovir. 5 concentration of ganciclovir required to inhibit 3Hthymidine incorporated by 50% was determined Additionally, a relative measure of adenovirus - mediated gene transfer and expression of each cell line was determined using a control virus which directs expression 10 of the marker gene beta-galactosidase. The data presented in Figure 14 and Table 1 below show that the ACNTK virus/ganciclovir combination treatment was capable of inhibiting cellular proliferation in all cell lines examined as compared with the control adenovirus ACN in 15 combination with ganciclovir. In contrast, the AANTK viral vector was only effective in those HCC cell lines which have been demonstrated to express α -fetoprotein. addition, the AANTK/GCV combination was more effective when the cells were plated at high densities.

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TABLE 1

| Cell Line | aFP | ß-gal Expression | ACN | ED50 ACNTK | AANTK |
|-------------|-----|---------------------|------|---------------|-------|
| MDAMB468 | + | +++ | >100 | 2 | >100 |
| BT549 | - | +++ | >100 | <0.3 | >100 |
| HLF | • | +++ | >100 | 0.8 | >100 |
| CHANG | • | +++ | >100 | 22 | >100 |
| HEP-3B | - | + | 80 | 8 | 8 |
| HEP-G2 LOW | + | ++ | 90 | 2 | 35 |
| HEP-G2 HIGH | + | ++ | 89 | 0.5 | 4 |

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Nude mice bearing Hep3B tumors (N=5/group) were treated intratumorally and peritumorally with equivalent

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doses of ACNTK or ACN control. Twenty-four hours after the first administration of recombinant adenovirus, daily treatment of ganciclovir was initiated in all mice. Tumor dimensions from each animal were measured twice weekly via calipers, and average tumor sizes are plotted in Figure 16. Average tumor size at day 58 was smaller in the ACNTK-treated animals but the difference did not reach statistical significance (p<0.09, unpaired t-test). These data support a specific effect of ACNTK on tumor growth in vivo. No significant differences in average body weight were detected between the groups.

Although the invention has been described with reference to the above embodiments, it should be understood that various modifications can be made without departing from the spirit of the invention. Accordingly, the invention is limited only by the claims that follow.

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What is claimed is:

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1. A recombinant adenovirus expression vector comprising a partial or total deletion of a protein IX DNA and a gene encoding a foreign protein.

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- 2. The recombinant adenovirus expression vector of claim 1, wherein the deletion of the protein IX gene sequence extends from about 3500 bp from the 5' viral termini to about 4000 bp from the 5' viral termini.
- 3. The recombinant adenovirus expression vector of claim 2 further comprising deletion of a non-essential DNA sequence in adenovirus early region 3 and/or early region 4.
- 4. The recombinant adenovirus expression vector of claim 2 further comprising deletion of a DNA sequences designated adenovirus Ela and Elb.
- 5. The recombinant adenovirus expression vector of claim 2 further comprising deletion of early region 3 and/or 4 and DNA sequences designated adenovirus Ela and Elb.
- 6. The recombinant adenovirus expression vector of claim 4 or 5 further comprising a deletion of up to forty nucleotides positioned 3' to the Ela and Elb and protein IX deletion and a foreign DNA molecule encoding a polyadenylation signal.
 - 7. The recombinant adenovirus expression vector of claims 1 to 6, wherein the adenovirus is a Group C adenovirus selected from a serotype 1, 2, 5 or 6.

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- 8. The recombinant adenovirus expression vector of claim 1, wherein the gene is a DNA molecule up to 2.6 kilobases.
- 9. The recombinant adenovirus expression vector of claim 6, wherein the gene is a DNA molecule up to 4.5 kilobases.
- 10. The recombinant adenovirus expression vector of claim 1, wherein the gene encodes a foreign functional protein or a biologically active fragment thereof.
- 11. The recombinant adenovirus expression vector of claim 10, wherein the gene encodes a foreign functional tumor suppressor protein or a biologically active fragment thereof.
- 12. The recombinant adenovirus expression vector of claim 1, wherein the gene encodes a suicide protein or functional equivalent thereof.
- 13. A transformed host cell comprising the recombinant adenovirus expression vector of claim 1 or 10.
- 14. The transformed host cell of claim 13, wherein the host cell is a procaryotic or eucaryotic cell.
- 15. A method for transforming a pathologic hyperproliferative mammalian cell comprising contacting the cell with the expression vector of claim 1.
- 16. A method of treating a pathology in an animal or mammal caused by the absence of a tumor suppressor gene or the presence of a pathologically mutated tumor suppressor gene comprising administering to the animal or mammal an effective amount of the vector of claim 1 containing a gene encoding a foreign functional protein

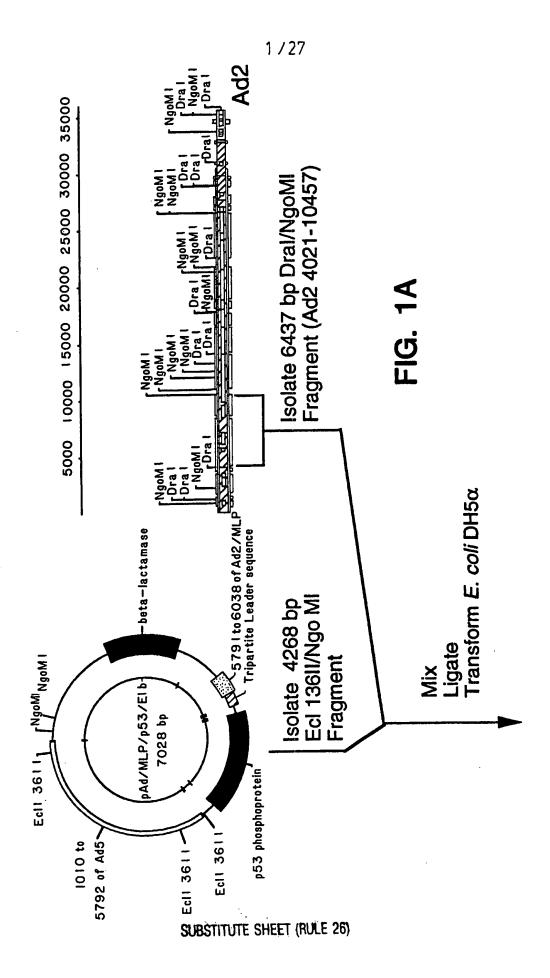
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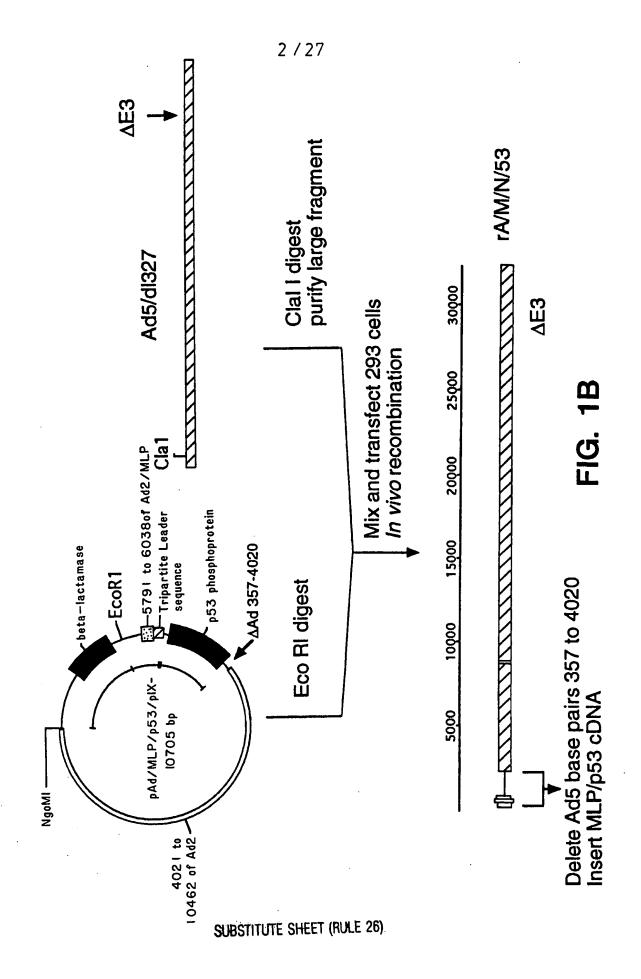
having a tumor suppressive function, under suitable conditions.

- 17. The method of claim 16, wherein the foreign protein is a functional tumor suppressor protein.
- 18. A method of gene therapy comprising administering to a subject an effective amount of the vector of claim 1.
- 19. A method of inhibiting the proliferation of a tumor in an animal comprising administering an effective amount of the adenoviral expression vector of claim 1 under suitable conditions to the animal.
- 20. The method of claim 19, wherein the gene encodes an anti-tumor agent.
- 21. The method of claim 20, wherein the antitumor agent is a tumor suppressor gene.
- 22. The method of claim 20, wherein the antitumor agent is a suicide gene or functional equivalent thereof.
- 23. The method of claim 21, wherein the tumor is non-small cell lung cancer, small cell lung cancer, hepatocarcinoma, melanoma, retinoblastoma, breast tumor, colorectal carcinoma, leukemia, lymphoma, brain tumor, cervical carcinoma, sarcoma, prostate tumor, bladder tumor, tumor of the reticuloendothelial tissues, Wilm's tumor, astrocytoma, glioblastoma, neuroblastoma, ovarian carcinoma, osteosarcoma, and renal cancer.
 - 24. The method of claim 19, wherein the vector is administered by intra-tumoral injection.

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- 25. A pharmaceutical composition comprising the recombinant adenoviral expression vector of claim 1, 10 or 12.
- 26. A method for reducing the proliferation of tumor cells in a subject comprising administering under suitable conditions an effective amount of an adenoviral expression vector of claim 12 and an effective amount of a thymidine kinase metabolite or a functional equivalent thereof.
 - 27. The method of claim 26, wherein the thymidine kinase metabolite is ganciclovir or 6-methoxypurine arabinonucleoside or a functional equivalent thereof.
 - 28. The method of claim 26, wherein the adenoviral expression vector is administered by injection into the tumor mass.
 - 29. The method of claim 26, wherein the tumor cells are hepatocellular carcinoma.
 - 30. The method of claim 29, wherein the adenoviral expression vector is administered directly into the hepatic artery of the subject.
- 31. A kit for reducing the proliferation of tumor cells comprising the components of the adenoviral expression vector of claim 12, a thymidine kinase metabolite or functional equivalent thereof, pharmaceutical carriers and instructions for the treatment of hepatocellular carcinoma using the kit components.





Met Pro Pro Lys Thr Pro Arg Lys Thr Ala Ala Thr Ala Ala Ala Ala Ala Ala Glu Pro Pro Ala Pro Pro Pro Pro Pro Pro Glu Glu Asp Pro Glu Gln Asp Ser Gly Pro Glu Asp Leu Pro Leu Val Arg Leu Glu Phe Glu Glu Thr Glu Glu Pro Asp Phe Thr Ala Leu Cys Gln Lys Leu Lys Ile Pro Asp His Val Arg Glu Arg Ala Trp Leu Thr Trp Glu Lys Val Ser Ser Val Asp Gly Val Leu Gly Gly Tyr Ile Gln Lys Lys Glu Leu Trp Gly Ile Cys Ile Phe Ile Ala Ala Val Asp Leu Asp Glu Met Ser Phe Thr Phe Thr Glu Leu Gln Lys Asn Ile Glu Ile Ser Val 115 His Lys Phe Phe Asn Leu Leu Lys Glu Ile Asp Thr Ser Thr Lys Val 135 Asp Asn Ala Met Ser Arg Leu Leu Lys Lys Tyr Asp Val Leu Phe Ala Leu Phe Ser Lys Leu Glu Arg Thr Cys Glu Leu Ile Tyr Leu Thr Gln 165 Pro Ser Ser Ser Ile Ser Thr Glu Ile Asn Ser Ala Leu Val Leu Lys 185 Val Ser Trp Ile Thr Phe Leu Leu Ala Lys Gly Glu Val Leu Gln Met 195 Glu Asp Asp Leu Val Ile Ser Phe Gln Leu Met Leu Cys Val Leu Asp 215 Tyr Phe Ile Lys Leu Ser Pro Pro Met Leu Leu Lys Glu Pro Tyr Lys 235 Thr Ala Val Ile Pro Ile Asn Gly Ser Pro Arg Thr Pro Arg Arg Gly 245 250

FIG. 2A

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Gln Asn Arg Ser Ala Arg Ile Ala Lys Gln Leu Glu Asn Asp Thr Arg 260 265 Ile Ile Glu Val Leu Cys Lys Glu His Glu Cys Asn Ile Asp Glu Val 280 Lys Asn Val Tyr Phe Lys Asn Phe Ile Pro Phe Met Asn Ser Leu Gly Leu Val Thr Ser Asn Gly Leu Pro Glu Val Glu Asn Leu Ser Lys Arg 310 Tyr Glu Glu Ile Tyr Leu Lys Asn Lys Asp Leu Asp Ala Arg Leu Phe 330 Leu Asp His Asp Lys Thr Leu Gln Thr Asp Ser Ile Asp Ser Phe Glu Thr Gln Arg Thr Pro Arg Lys Ser Asn Leu Asp Glu Glu Val Asn Val 360 Ile Pro Pro His Thr Pro Val Arg Thr Val Met Asn Thr Ile Gln Gln 370 375 Leu Met Met Ile Leu Asn Ser Ala Ser Asp Gln Pro Ser Glu Asn Leu 390 395 Ile Ser Tyr Phe Asn Asn Cys Thr Val Asn Pro Lys Glu Ser Ile Leu 410 Lys Arg Val Lys Asp Ile Gly Tyr Ile Phe Lys Glu Lys Phe Ala Lys Ala Val Gly Gln Gly Cys Val Glu Ile Gly Ser Gln Arg Tyr Lys Leu Gly Val Arg Leu Tyr Tyr Arg Val Met Glu Ser Met Leu Lys Ser Glu 450 455 Glu Glu Arg Leu Ser Ile Gln Asn Phe Ser Lys Leu Leu Asn Asp Asn 470 Ile Phe His Met Ser Leu Leu Ala Cys Ala Leu Glu Val Val Met Ala 490 Thr Tyr Ser Arg Ser Thr Ser Gln Asn Leu Asp Ser Gly Thr Asp Leu 505 500

FIG. 2B

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Ser Phe Pro Trp Ile Leu Asn Val Leu Asn Leu Lys Ala Phe Asp Phe Tyr Lys Val Ile Glu Ser Phe Ile Lys Ala Glu Gly Asn Leu Thr Arg Glu Met Ile Lys His Leu Glu Arg Cys Glu His Arg Ile Met Glu Ser Leu Ala Trp Leu Ser Asp Ser Pro Leu Phe Asp Leu Ile Lys Gln Ser 570 Lys Asp Arg Glu Gly Pro Thr Asp His Leu Glu Ser Ala Cys Pro Leu Asn Leu Pro Leu Gln Asn Asn His Thr Ala Ala Asp Met Tyr Leu Ser Pro Val Arg Ser Pro Lys Lys Gly Ser Thr Thr Arg Val Asn Ser Thr Ala Asn Ala Glu Thr Gln Ala Thr Ser Ala Phe Gln Thr Gln Lys 630 640 Pro Leu Lys ser Thr Ser Leu Ser Leu Phe Tyr Lys Lys Val Tyr Arg 650 Leu Ala Tyr Leu Arg Leu Asn Thr Leu Cys Glu Arg Leu Leu Ser Glu His Pro Glu Leu Glu His Ile Ile Trp Thr Leu Phe Gln His Thr Leu 680 Gln Asn Glu Tyr Glu Leu Met Arg Asp Arg His Leu Asp Gln Ile Met 690 Met Cys Ser Met Tyr Gly Ile Cys Lys Val Lys Asn Ile Asp Leu Lys Phe Lys Ile Ile Val Thr Ala Tyr Lys Asp Leu Pro His Ala Val Gln Glu Thr Phe Lys Arg Val Leu Ile Lys Glu Glu Glu Tyr Asp Ser Ile 740 Ile Val Phe Tyr Asn Ser Val Phe Met Gln Arg Leu Lys Thr Asn Ile 760

FIG. 2C

Leu Gln Tyr Ala Ser Thr Arg Pro Pro Thr Leu Ser Pro Ile Pro His 770 775 780 Ile Pro Arg Ser Pro Tyr Lys Phe Pro Ser Ser Pro Leu Arg Ile Pro 795 Gly Gly Asn Ile Tyr Ile Ser Pro Leu Lys Ser Pro Tyr Lys Ile Ser 805 Glu Gly Leu Pro Thr Pro Thr Lys Met Thr Pro Arg Ser Arg Ile Leu Val Ser Ile Gly Glu Ser Phe Gly Thr Ser Glu Lys Phe Gln Lys Ile Asn Gln Met Val Cys Asn Ser Asp Arg Val Leu Lys Arg Ser Ala Glu 850 Gly Ser Asn Pro Pro Lys Pro Leu Lys Lys Leu Arg Phe Asp Ile Glu 875 880 Gly Ser Asp Glu Ala Asp Gly Ser Lys His Leu Pro Gly Glu Ser Lys Phe Gln Gln Lys Leu Ala Glu Met Thr Ser Thr Arg Thr Arg Met Gln 910 Lys Gln Lys Met Asn Asp Ser Met Asp Thr Ser Asn Lys Glu Glu Lys 915 920

FIG. 2D

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| TTC | GGT: | TTT | TCTC | AGGGG | GA C | g tt G/ | TAAA | TA 1 | rttt(| STAA | CGG | GAGT | CGG | GAGA | GGACGG | 6 |
|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-----|
| GGC | TGC | ccc | GCGT | GCGC | GC G | CGTC | STCCI | CC | CCGG | CGCT | CCT | CCAC | AGC | TCGC | TGGCTC | 12 |
| CCG | CCCC | GA | AAGG | CGTC | | | CCC Pro | | | | | | | | | 17: |
| ACC Thr | GCC Ala | GCC Ala | GCT Ala 15 | GCC Ala | GCC Ala | GCG Ala | GAA Glu | CCC Pro 20 | CCG Pro | GCA Ala | CCG Pro | CCG Pro | CCG Pro 25 | CCG Pro | CCC Pro | 219 |
| CCT Pro | CCT Pro | GAG Glu 30 | GAG Glu | GAC Asp, | CCA Pro | GAG Glu | CAG Gln 35 | GAC Asp | AGC Ser | GGC Gly | CCG Pro | GAG Glu 40 | GAC Asp | CTG Leu | CCT Pro | 267 |
| CTC Leu | GTC Val 45 | AGG Arg | CTT | GAG Glu | TTT Phe | GAA Glu 50 | GAA Glu | ACA Thr | GAA Glu | GAA Glu | CCT Pro 55 | GAT Asp | TTT Phe | ACT Thr | GCA Ala | 315 |
| TTA Leu 60 | TGT Cys | CAG Gln | AAA Lys | TTA Leu | AAG Lys 65 | ATA Ile | CCA Pro | GAT Asp | CAT His | GTC Val 70 | AGA Arg | GAG Glu | AGA Arg | GCT Ala | TGG Trp 75 | 363 |
| TTA Leu | ACT Thr | TGG Trp | GAG Glu | AAA Lys 80 | GTT Val | TCA Ser | TCT Ser | GTG Val | GAT Asp 85 | GGA Gly | GTA Val | TTG Leu | GGA Gly | GGT Gly 90 | TAT Tyr | 411 |
| ATT Ile | CAA Gln | AAG Lys | AAA Lys 95 | AAG Lys | GAA Glu | CTG Leu | TGG Trp | GGA Gly 100 | ATC Ile | TGT Cys | ATC Ile | TTT Phe | ATT Ile 105 | GCA Ala | GCA Ala | 459 |
| GTT Val | GAC Asp | CTA Leu 110 | GAT Asp | GAG Glu | ATG Met | TCG Ser | TTC Phe 115 | ACT Thr | TTT Phe | ACT Thr | GAG Glu | CTA Leu 120 | CAG Gln | AAA Lys | AAC Asn | 507 |
| ATA Ile | GAA Glu 125 | ATC | AGT Ser | GTC Val | CAT His | AAA Lys 130 | TTC Phe | TTT Phe | AAC Asn | TTA Leu | CTA Leu 135 | AAA Lys | GAA Glu | ATT | GAT Asp | 555 |
| ACC Thr 140 | AGT Ser | ACC Thr | AAA Lys | GTT Val | GAT Asp 145 | AAT Asn | GCT Ala | ATG Met | TCA Ser | AGA Arg 150 | CTG Leu | TTG Leu | AAG Lys | AAG Lys | TAT Tyr 155 | 603 |
| GAT Asp | GTA Val | TTG Leu | TTT | GCA Ala 160 | CTC Leu | TTC Phe | AGC Ser | AAA Lys | TTG Leu 165 | GAA Glu | AGG Arg | ACA Thr | TGT Cys | GAA Glu 170 | CTT Leu | 651 |
| ATA Ile | TAT Tyr | TTG Leu | ACA Thr 175 | CAA Gln | CCC Pro | AGC Ser | AGT Ser | TCG Ser 180 | ATA Ile | TCT Ser | ACT Thr | GAA Glu | ATA Ile 185 | AAT Asn | TCT Ser | 699 |
| GCA Ala | TTG Leu | GTG Val 190 | CTA Leu | AAA Lys | GTT Val | TCT Ser | TGG Trp 195 | ATC Ile | ACA Thr | TTT Phe | TTA Leu | TTA Leu 200 | GCT Ala | AAA Lys | GGG Gly | 747 |
| GAA Glu | GTA Val | Leu | CAA Gln | ATG Met | GAA Glu | GAT Asp | GAT Asp | CTG Leu | GTG Val | ATT Ile | TCA Ser | TTT Phe | CAG Gln | TTA Leu | ATG Met | 795 |

FIG. 3A

SUBSTITUTE SHEET (RULE 26)

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| | | | | | | | • | | | | | | | | | |
|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|------------|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| | TGT Cys | | | | | | | | | | | | | | | 843 |
| | GAA Glu | | | | | | | | | | | | | | | 891 |
| | CCC Pro | | | | | | | | | | | | | | | 939 |
| | AAT Asn | | | | | | | | | | | | | | | 987 |
| | ATA Ile 285 | | | | | | | | | | | | | | | 1035 |
| | AAT Asn | | | | | | | | | | | | | | | 1083 |
| AAT | CTT Leu | TCT Ser | AAA Lys | CGA Arg 320 | TAC Tyr | GAA Glu | GAA Glu | ATT Ile | TAT Tyr 325 | CTT Leu | AAA Lys | AAT Asn | AAA Lys | GAT Asp 330 | CTA Leu | 1131 |
| GAT Asp | GCA Ala | AGA Arg | TTA Leu 335 | TTT Phe | TTG Leu | GAT Asp | CAT His | GAT Asp 340 | AAA Lys | ACT Thr | CTT Leu | CAG Gln | ACT Thr 345 | GAT Asp | TCT Ser | 1179 |
| ATA Ile | GAC Asp | AGT Ser 350 | TTT Phe | GAA Glu | ACA Thr | CAG Gln | AGA Arg 355 | ACA Thr | CCA Pro | CGA Arg | AAA Lys | AGT Ser 360 | AAC Asn | CTT Leu | GAT Asp | 1227 |
| GAA Glu | GAG Glu 365 | GTG Val | TAA Asn | GTA Val | ATT | CCT Pro 370 | CCA Pro | CAC His | ACT Thr | CCA Pro | GTT Val 375 | AGG Arg | ACT Thr | GTT Val | ATG Met | 1275 |
| AAC Asn 380 | ACT Thr | ATC Ile | CAA Gln | CAA Gln | TTA Leu 385 | Met | ATG Met | Ile | Leu | Asn | TCA Ser | GCA Ala | AGT Ser | GAT Asp | CAA Gln 395 | 1323 |
| CCT Pro | TCA Ser | GAA Glu | AAT Asn | CTG Leu 400 | ATT Ile | TCC Ser | TAT Tyr | TTT Phe | AAC Asn 405 | AAC Asn | TGC Cya | ACA Thr | GTG Val | AAT Asn 410 | CCA Pro | 1371 |
| AAA Lys | GAA Glu | AGT Ser | ATA Ile 415 | CTG Leu | AAA Lys | AGA Arg | GTG Val | AAG Lys 420 | GAT Asp | ATA Ile | GGA Gly | TAC Tyr | ATC Ile 425 | TTT Phe | AAA Lys | 1419 |
| GAG Glu | Lys | TTT Phe 430 | GCT Ala | AAA Lys | GCT Ala | GTG Val | GGA Gly 435 | CAG Gln | GGT Gly | TGT Cys | GTC Val | GAA Glu 440 | ATT | GGA Gly | TCA Ser | 1467 |
| CAG Glr | CGA Arg 445 | TAC Tyr | ГЛа УУУ | CTT Leu | GGA Gly | GTT Val 450 | CGC Arg | TTG Leu | TAT Tyr | TAC Tyr | CGA Arg 455 | GTA Val | ATG Met | GAA Glu | TCC Ser | 1515 |

FIG. 3B

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| | | | | | | | | TTA Leu | | | | | | | | 1563 |
|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| | | | | | | | | ATG Met | | | | | | | | 1611 |
| | | | | | | | | AGA Arg 500 | | | | | | | | 1659 |
| | | | | | | | | TGG Trp | | | | | | | | 1707 |
| AAA Lys | GCC Ala 525 | TTT Phe | GAT Asp | TTT Phe | TAC Tyr | AAA Lys 530 | GTG Val | ATC Ile | GAA Glu | AGT Ser | TTT Phe 535 | ATC Ile | AAA Lys | GCA Ala | GAA Glu | 1755 |
| GGC Gly 540 | AAC Asn | TTG Leu | ACA Thr | AGA Arg | GAA Glu 545 | ATG Met | ATA Ile | AAA Lys | CAT His | TTA Leu 550 | GAA Glu | CGA Arg | TGT Cys | GAA Glu | CAT His 555 | 1803 |
| CGA Arg | ATC Ile | ATG Met | GAA Glu | TCC ser 560 | CTT Leu | GCA Ala | TGG Trp | CTC Leu | TCA Ser 565 | GAT Asp | TCA Ser | CCT Pro | TTA Leu | TTT Phe 570 | GAT Asp | 1851 |
| CTT Leu | ATT Ile | AAA Lys | CAA Gln 575 | TCA Ser | AAG Lys | GAC Asp | CGA Arg | GAA Glu 580 | GGA Gly | CCA Pro | ACT Thr | GAT Asp | CAC His 585 | CTT Leu | GAA Glu | 1899 |
| TCT | GCT Ala | TGT Cya 590 | CCT Pro | CTT Leu | AAT Asn | CTT Leu | CCT Pro 595 | CTC Leu | CAG Gln | TAA Asn | TAA neA | CAC His 600 | ACT Thr | GCA Ala | GCA Ala | 1947 |
| GAT Asp | ATG Met 605 | TAT Tyr | CTT Leu | TCT Ser | CCT Pro | GTA Val 610 | AGA Arg | TCT Ser | CCA Pro | AAG Lys | AAA Lys 615 | AAA Lys | GGT Gly | TCA Ser | ACT Thr | 1995 |
| ACG Thr 620 | CGT Arg | GTA Val | AAT Asn | TCT Ser | ACT Thr 625 | GCA Ala | TAA neA | GCA Ala | GAG Glu | ACA Thr 630 | CAA Gln | GCA Ala | ACC Thr | TCA Ser | GCC Ala 635 | 2043 |
| TTC Phe | CAG Gln | ACC Thr | CAG Gln | AAG Lys 640 | CCA Pro | TTG Leu | AAA Lys | TCT Ser | ACC Thr 645 | TCT Ser | CTT Leu | TCA Ser | CTG Leu | TTT Phe 650 | TAT Tyr | 2091 |
| AAA Lys | AAA Lys | GTG Val | TAT Tyr 655 | CGG Arg | CTA Leu | GCC Ala | TAT Tyr | CTC Leu 660 | CGG Arg | CTA Leu | TAA naA | ACA Thr | CTT Leu 665 | TGT Cys | GAA Glu | 2139 |
| CGC Arg | CTT | CTG Leu 670 | TCT | GAG Glu | CAC His | CCA Pro | GAA Glu 675 | TTA Leu | GAA Glu | CAT His | ATC Ile | ATC Ile 680 | TGG Trp | ACC Thr | CTT Leu | 2187 |
| TTC Phe | CAG Gln 685 | CAC His | ACC Thr | CTG Leu | CAG Gln | AAT Asn 690 | GAG Glu | TAT Tyr | GAA Glu | CTC Leu | ATG Met 695 | AGA Arg | GAC Asp | AGG Arg | CAT His | 2235 |

FIG. 3C

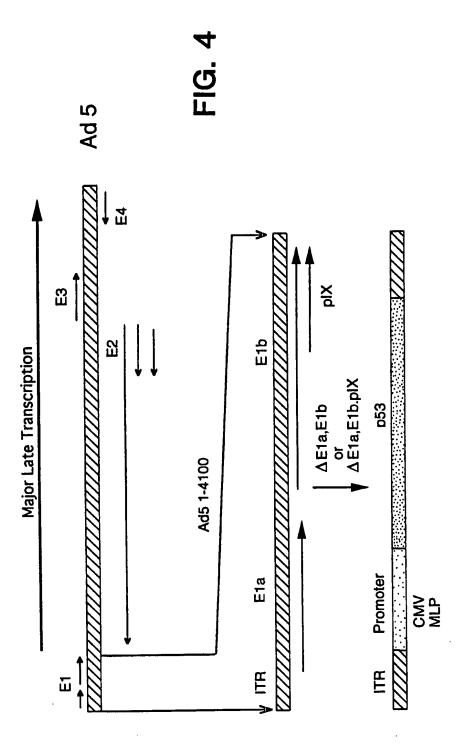
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| | | | | 107 | _ / | | | | | | | | |
|--|------------|-----|-------|-----|-------------------|------|-----|-------|-------|-------|-------|------------|------|
| | | | | | ATG Met | | | | | | | | 2283 |
| | | | | | ATT Ile | | | | | | | | 2331 |
| | | | | | AAA Lys 740 | | | | | | | GAG Glu | 2379 |
| | | | | | TAT Tyr | | | | | | | | 2427 |
| | | | | | GCT Ala | | | | | | | | 2475 |
| | | | | | AGC Ser | | | | | | | | 2523 |
| | | | | | ATC Ile | | | | | | | | 2571 |
| | | | | | CCA Pro 820 | | | | | | | | 2619 |
| | | | | | GGT Gly | | | | | | | | 2667 |
| | | | | | GTA Val | | | | | | | | 2715 |
| | | | | | CCT Pro | | | | | | | | 2763 |
| | | | | | GAA Glu | | | | | | | | 2811 |
| | | | | | AAA Lys 900 | | | | | | | | 2859 |
| | | | | | ATG Met | | | | | | | | 2907 |
| | GAG Glu | TGA | GGAT(| CTC | AGGA | CCTT | G T | GGAC: | ACTG: | r GT1 | ACACO | CTCT | 2962 |

GGATTCATTG TCTCTCACAG ATGTGACTGA TAT

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FIG. 3D SUBSTITUTE SHEET (RULE 26)



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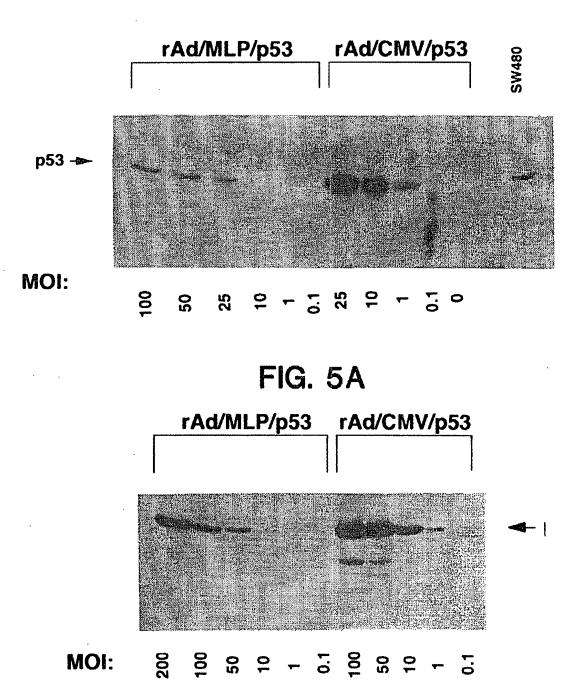


FIG. 5B

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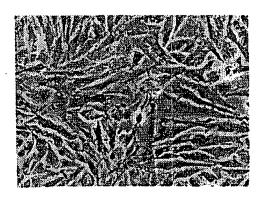


FIG. 6A

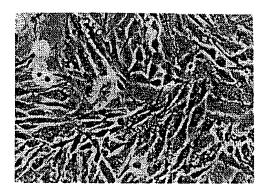


FIG. 6B

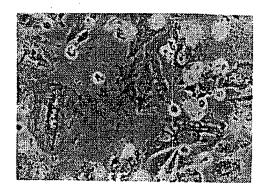
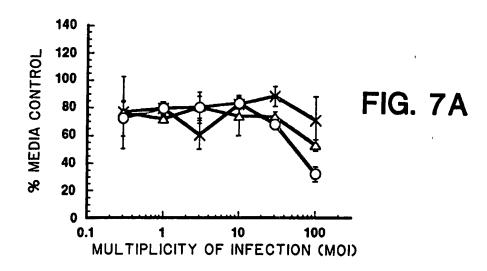
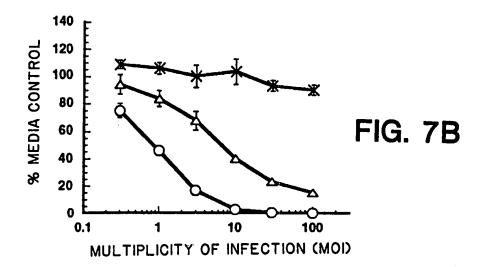
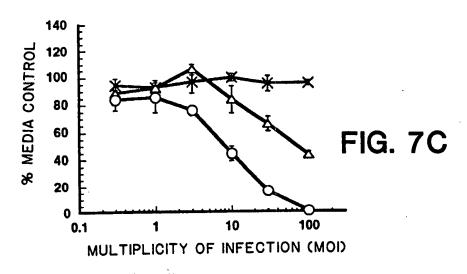


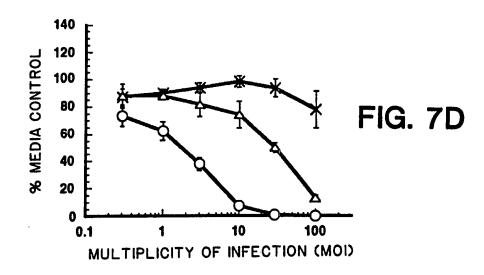
FIG. 6C

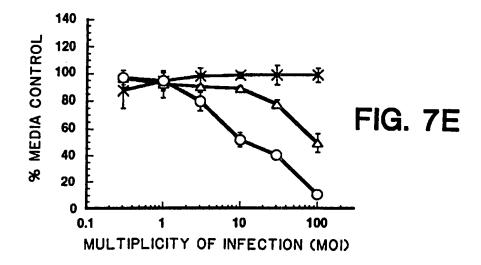


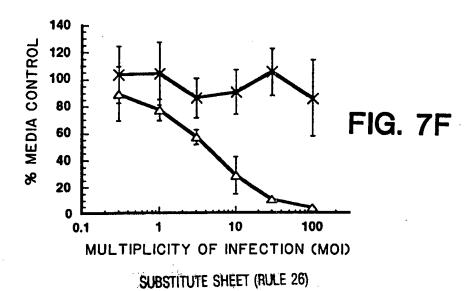




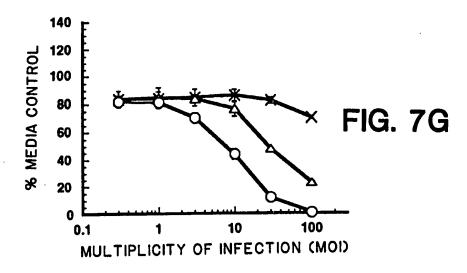
SUBSTITUTE SHEET (RULE 26)

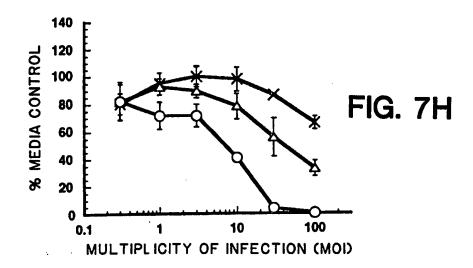


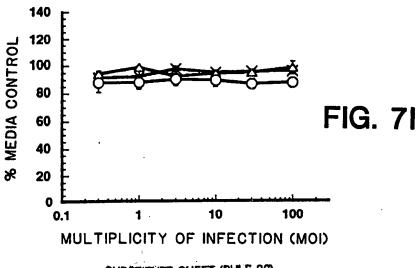




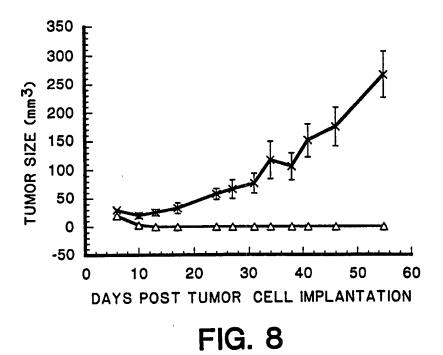








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SUBSTITUTE SHEET (RULE 26)

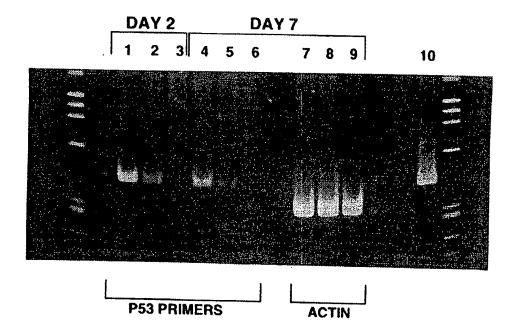


FIG. 9

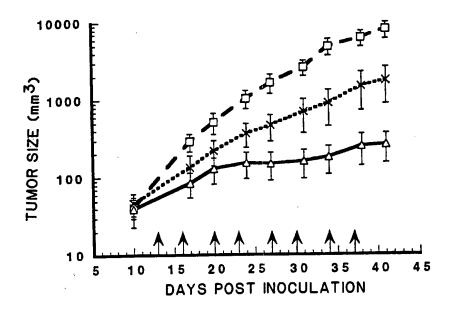


FIG. 10A

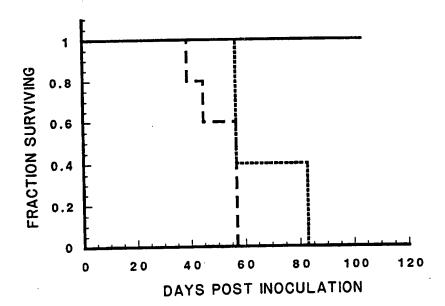
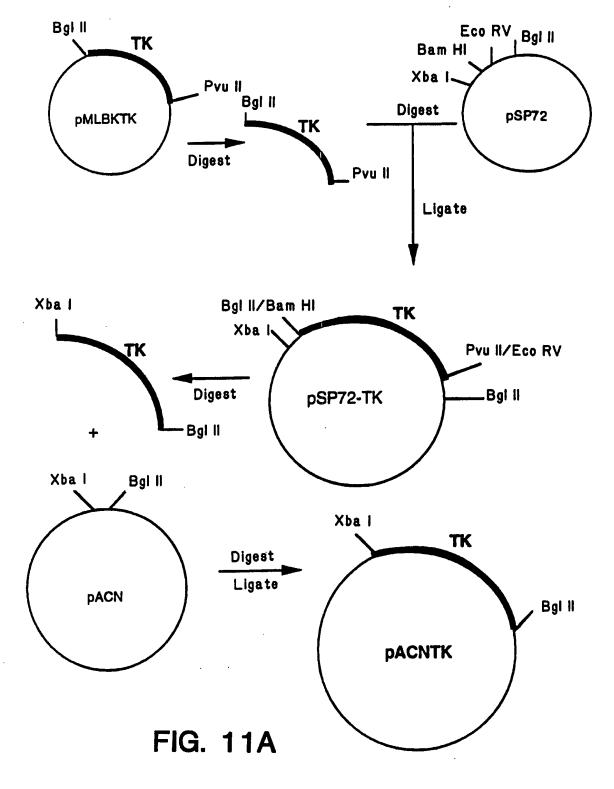
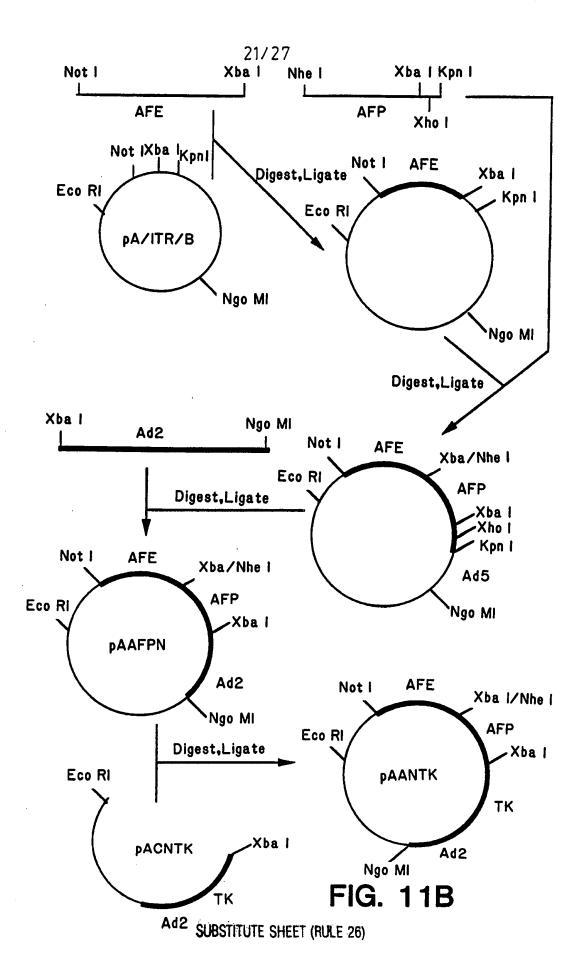


FIG. 10B

20/27





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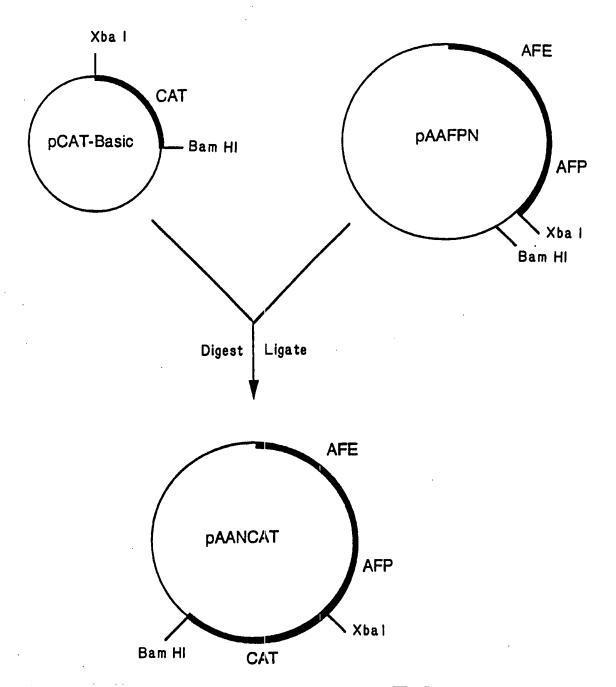
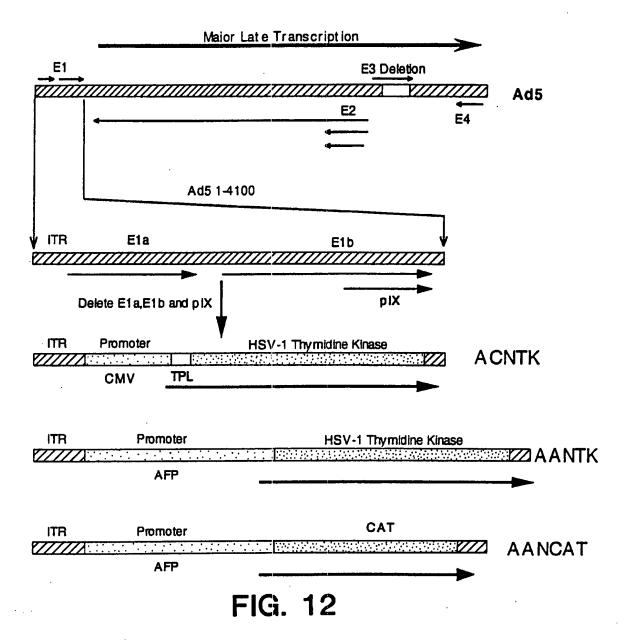
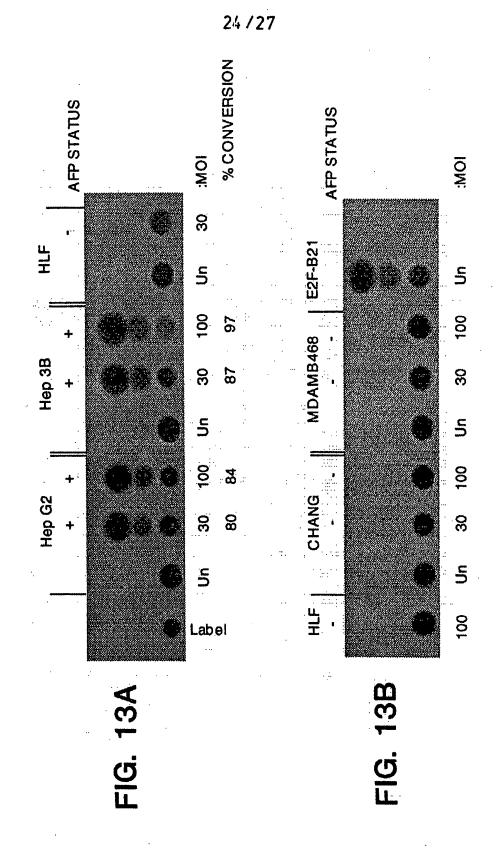
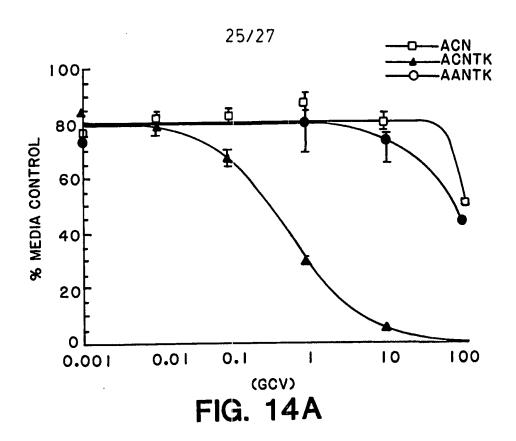


FIG. 11C



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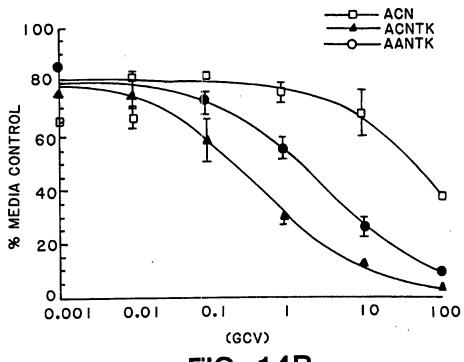


FIG. 14B SUBSTITUTE SHEET (RULE 26)

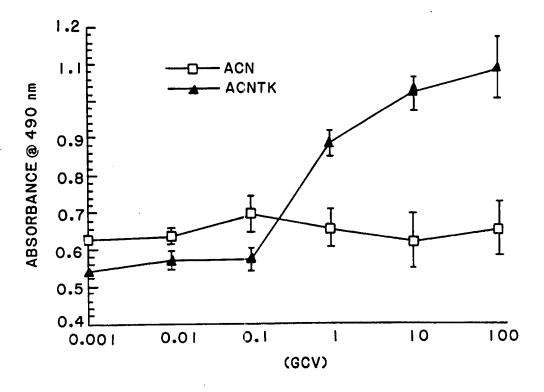


FIG. 15

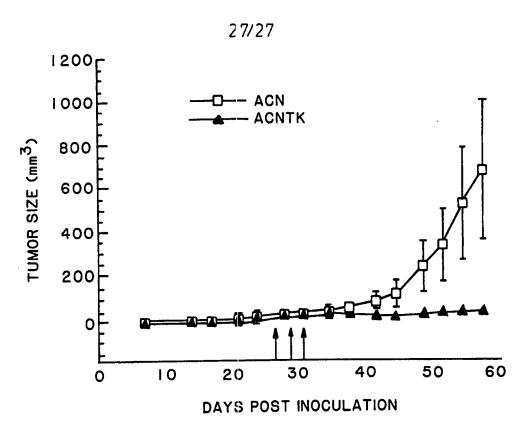


FIG. 16A

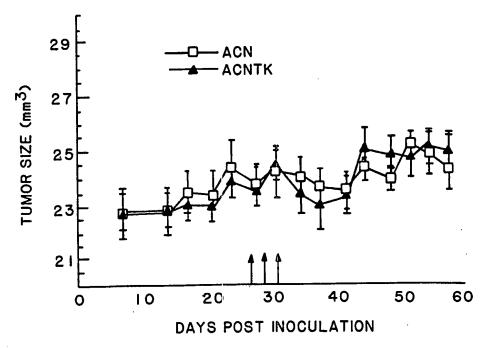


FIG. 16B

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IDS REFERENCES

FOR

EPO - Munich 45 **1 7. Sep. 2007**



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September 14, 2007

Publication No.:

0 833 934

Application No.:

96 917 735.1

Appeal No.:

T 1205/07

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Our Ref.:

OP33

OPPONENT'S STATEMENT OF GROUNDS OF APPEAL

This constitutes the written statement setting out the grounds of appeal of the Appellant (Opponent 01), Merck Serono International S.A., against the decision of the Opposition Division (OD) dated May 11, 2007 maintaining European Patent No. 0 833 934 in amended form.

The OD decided that the claims of the patent as granted contravened both Art 100(c) and Art. 100(b). We have no quarrel with this part of the decision. However, the OD also decided that AR2, filed by the Patentee during the oral proceedings to overcome the above-mentioned deficiencies, satisfied all requirements under the EPC, including those of Art. 100(a) EPC. The opposed patent was thus maintained in amended form based on the claims of AR2.

Appellant hereby requests that the opposed patent be revoked in its entirety based on novelty (Art. 100(a) and 54 EPC) and of lack of inventive step (Art. 100(a) and 56 EPC)

It is further requested to admit further documents as specified below under item 2 into the proceedings.

An oral hearing is requested if the Board of Appeal does not intend to revoke the patent in the light of the written submissions.

1. New documents filed with the present submission

The opponent submits herewith the four (4) following documents, which will be referred to by the numbers given below.

Documents D52 to D55 are highly relevant to the validity of the opposed patent.

D52 Figure 2 by Opponent 01.

This Figure illustrates the inventive step argument made in section 5.1.1. of this submission.

D53 van den Elsen et al. (1982) "The relationship between region E1a and E1b of human adenoviruses in cell transformation" Gene. 18(2):175-85.

This article discloses a BRK cell line transformed with a fragment consisting of Ad5 nucleotides 1045-3978, i.e. lacking pIX sequences. It is thus highly relevant for inventive step.

D54 Kawarabayasi and Sugisaki (1985) "Structure of viral DNA in a rat cell line transformed by the cloned EcoRI-C fragment of adenovirus 12." Nucleic Acids Res. 13(18):6591-604.

This article discloses a rat cell line transformed with a fragment consisting of Ad12 nucleotides 118-3520, i.e. lacking pIX sequences. It is thus highly relevant for inventive step.

D55 Kimura et al. (1981) "Nucleotide sequence of the transforming early region E1b of adenovirus type 12 DNA: structure and gene organization, and comparison with those of adenovirus type 5 DNA." Nucleic Acids Res. 9(23):6571-89.

This article compares the sequence of the E1 regions of Ad5 and of Ad12, and is thus useful for analyzing D54.

2. The claims of the patent as maintained in amended form

In addition to the cancellation of claims 7, 9, 12-19, 21, 22, 24-26, 29, 31, 33 and 36, there are three main differences between the claims of the patent as granted and the claims as maintained in amended form (AR2). More specifically, in AR2:

- the packaging cell is defined as a cell comprising in its genome an adenoviral sequence consisting of Ad5 nucleotides 459-3510 (vs. a cell that encodes the adenoviral E1A and E1B proteins but lacks pIX sequences in the patent as granted);
- the cells are limited to Human Embryonic Retinoblast (HER), Human Embryonic Kidney (HEK), or a Human Embryonic Lung (HEL) cells; and

- the term "packaging system" is replaced with the term "combination".

The broadest claim of the patent as maintained in amended form is claim 19, which is directed to a packaging cell characterized in that it comprises in its genome an adenoviral sequence consisting of Ad5 nucleotides 459-3510, and wherein said cell is a HER, HEK or HEL cell.

Claim 1 is directed to a combination of the packaging cell of claim 19 with an adenoviral vector, wherein said adenoviral vector has no overlapping sequences which allow for homologous recombination between adenovirus DNA present in the packaging cell and the vector.

Claims 2, 3, 5-9, 11-13, 15 and 18 all depend either directly or indirectly on claim 1.

Claim 4 is directed to a combination of a packaging cell with an adenoviral vector, wherein said combination comprises an adenoviral vector resulting from the action of a nucleic acid polymerase comprised by the combination of claim 3.

Claim 10 is redundant with claim 19.

Claim 16 is directed to a specific cell according to claim 10 that has been deposited at the ECACC.

Claim 17 is directed to a packaging cell comprising in its genome an adenoviral sequence encoding at least both the E1A and a mutated E2A region, wherein at least one of the E2A products is temperature sensitive, and wherein said cell is a HER, HEK or HEL cell.

3. Novelty (Art. 100(a) & Art 54 EPC)

Claim 17 of AR2 is not novel over D22. The Board is directed to section 5.4. of the Grounds of Opposition filed by Opponent 02.

During the oral proceedings, the limitation "wherein said cell is a HER, HEK or HEL cell" was added to original claim 35. However, this amendment is not sufficient to gain novelty over D22. Specifically, in D22 the Ad2ts111Sal1A and Ad2EcoRIA packaging constructs were transfected into the 293 cell line, thereby providing the 293 cell line with packaging nucleic acids that provide the transfected cells with the ability to produce adenoviral gene products from the E1A and E2 regions, wherein the packaging nucleic acid molecule encoding the E2A region is mutated so that at least one E2A product is temperature sensitive (see e.g. page 886, left column, lines 11-15 of D22). As evidenced by D4, the 293 cell line is a HEK cell line (see e.g. first sentence of the abstract of D4).

Therefore, claim 17 lacks novelty over D22.

4. Inventive Step (Art. 100(a) & Art 56 EPC)

The OD decided that all claims of AR2 were allegedly inventive. Contrary to the OD's finding, it will be demonstrated herebelow that claims 1-16, 18 and 19 of AR2 lack inventive step.

We will first deal with claim 19, which is the broadest claim of the patent according to AR2.

4.1. Claim 19 lacks inventive step

In the proceedings, Opponent 01 had combined D21 with the knowledge of the person skilled in the art illustrated by e.g. D3 to arrive at the cell of claim 19 in an obvious manner (see section e.g. 6.6.1. of the decision). We maintain that this is the case.

4.1.1. Problem solution approach

(i) D21 as closest prior art

D21, published in December 1994, provides both adenoviral vectors and packaging cell lines allowing to propagate such vectors (see e.g. title of D21). In addition, D21 aims at overcoming the problem lying in the generation of unwanted replication-competent adenoviruses (RCA), as indicated in column 3, lines 24-27 of D21a, the description of which corresponds to an English translation of D21. D21 is thus considered to be the closest prior art.

(ii) Difference between claim 19 and D21

Claim 19 is directed to a packaging cell characterized in that it comprises in its genome an adenoviral sequence consisting of Ad5 nucleotides 459-3510, and wherein said cell is a Human Embryonic Retinoblast (HER) cell, a Human Embryonic Kidney (HEK), or a Human Embryonic Lung (HEL) cell. The E1 region of the Ad5 genome was well-known in the art at the priority date of the opposed patent (see e.g. column 13, lines 64-65 of D21a). The person skilled in the art thus knew that nucleotides 459-3510 of Ad5 comprised the sequences coding for the E1A proteins, for E1B-21kD, and for E1B-55kD.

D21 discloses a complementation cell comprising a fragment of the Ad5 genome (see column 10, lines 53-57 of D21a). This complementation cell is used for packaging adenoviruses (see title of D21a). As indicated in column 10, lines 4-7 of D21a, the complementation cell of D21 encodes the E1A proteins, E1B-21kD, and E1B-55kD. In addition, the complementation cell may be a HER cell (see e.g. column 12, lines 6-8 of D21a).

Thus, the only difference between the packaging cell of claim 19 and the packaging cell of D21 lies in the boundaries of the specific fragment integrated in the genome of the cells, i.e. nucleotide 459 at the 5' end and nucleotide 3510 at the 3' end.

(iii) Objective technical problem & solution provided by the opposed patent

The objective technical problem to be solved by claim 19 of the opposed patent is therefore the provision of an alternative packaging cell line.

The solution of claim 19 of the opposed patent is a HER, HEK or HEL cell comprising in its genome nucleotides 459-3510 of Ad5.

It will be shown herebelow that this solution was obvious in view of D21 combined with the knowledge of the person skilled in the art. Figure 2 (D52) enclosed herewith is submitted to illustrate our argument.

(iv) This solution was obvious in view of the prior art

D21 pertains to a vector deleted for all or part of the E1A region and the whole of the sequences coding for the early proteins of the E1B region (column 4, lines 4-6 of D21a).

In the embodiment wherein all the E1A region is deleted, the 5' boundary of the deletion must be located around nucleotide 468 of Ad5, which the person skilled in the art knew to be the position of the E1A promoter (see e.g. page 3, line 33 of D3). Thus, D21 teaches a vector in which the 5' boundary is located around nucleotide 468 of Ad5.

Additional information about the 3' boundary of such a deletion is given in the same paragraph (see column 4, lines 42-45 of D21a). It is specifically indicated that the deletion must necessarily encompass nucleotides 1634-3509 of the Ad5 E1 region. Accordingly, D21 teaches a vector in which the 3' boundary is located at nucleotide 3509 of Ad5.

Thus this paragraph of D21 suggests a vector deleted for nucleotide 468-3509 of Ad5.

Since the packaging cell line must provide all the functions deleted in the vector (see column 1, lines 5-10 of D21a), and since one should eliminate the sequences in common to the vector and the Ad fragment integrated into the packaging cell (see column 4, lines 42-51 of D21a), the person skilled in the art knows that the packaging cell line must provide the functions missing in the vector but nothing more. As a matter of fact, the person skilled in the art would see the packaging line as a mirror image of the vector, said packaging cell providing the functions missing in the vector. Hence, the packaging line is directly and unambiguously derivable from the vector by the person skilled in the art.

In the present case, the mirror image of an Ad5 vector deleted in nucleotides 468-3509 is a packaging cell line comprising in its genome nucleotides 468-3509 of Ad5.

As apparent from D3, an Ad5 fragment consisting of nucleotides 459-3510 encodes exactly the same functions as a fragment consisting of nucleotides 468-3509, i.e. the E1A and E1B early proteins. On the other hand, pIX is missing in either Ad5 fragment. As also apparent from D3, no function has been assigned to nucleotides 459 to 468 of Ad5. Therefore, the presence of these nine additional nucleotides in the packaging cell line of claim 19 cannot have any technical effect.

Thus, the packaging cell of claim 19 of the opposed patent is a mere equivalent without technical effect of the packaging cell line suggested in D21. In conclusion, the packaging cell of claim 19 lacks inventive step in view of D21 combined with the knowledge of the person skilled in the art.

4.1.2. The OD's decision

In section 6.6.2 of the decision the OD sets out the reasons why it came to the conclusion of the existence of an inventive step. We will now discuss the OD's arguments in detail, and show that the OD's reasoning is flawed.

(i) Omission of pIX in the packaging cell line

The OD asserts that there is no teaching in D21 to omit pIX from the packaging cell, neither implicit and certainly not explicit. The OD cites Example 2 and the passage in column 10, lines 58-65 to support his assertion.

However, the OD has disregarded the passage in column 10, lines 4-7 of D21a, which provides an unambiguous implicit support for a packaging cell lacking pIX sequences:

"According to an advantageous embodiment, a complementation line according to the invention comprises, in particular, all or part of the E1A region and the whole of the sequences coding for the <u>early proteins of the E1B region</u>."

D21 thus teaches a preferred packaging cell line that comprises the sequences coding for E1A and the <u>early</u> proteins of the E1B region.

The fact that this packaging cell line has been defined as "comprising sequences coding for the early proteins of the E1B region" and <u>not</u> as "comprising the E1B region" must be interpreted as a teaching to use E1B without pIX sequences. This wording necessarily excludes pIX, which was known to be a late protein (see e.g. column 4, line 42 of D21a).

This specific wording cannot be seen as an arbitrary choice of the authors of D21. As a matter of fact, D21 constantly makes a difference between the early proteins of the E1B region and the late pIX protein.

In the "Background of the Invention" section, D21 differentiates the E1B proteins from the pIX protein, and teaches that there is an overlap between the transcription termination signal of the E1B proteins and the sequence coding for the pIX protein:

"The gene coding for structural protein IX of which the capsid is composed is located at the 5' end of the adenoviral genome and overlaps the E1B region at its 3' end. The protein IX transcription unit utilizes the same transcription termination signal as the E1B transcription unit."

(column 2, lines 36-41 of D21a)

Later on, when describing the adenoviral vectors to be used with the packaging cell lines, D21a further clarifies in column 4, lines 32-42 what is meant by the "sequences coding for the E1B early proteins":

"According to a first variant, an adenoviral vector according to the invention is derived from the genome of a natural or wild-type adenovirus by deletion of all or part of the E1A region and the portion of the E1B region comprising the whole of the sequences coding for the early proteins. According to a preferred embodiment, the deletion affects the promoter and the sequences coding for the expression products of the E1B region, that is to say the early proteins, and does not include all or part of the transcription termination signal which overlaps the sequences coding for the late protein IX."

Therefore, the sequences coding for the E1B early proteins do <u>not</u> include the transcription termination signal of the E1B early proteins, which overlaps the sequences coding for the late protein IX. In other words, D21 unambiguously teaches that the sequences coding for the E1B <u>early</u> proteins do <u>not</u> comprise any pIX sequences.

In view of the above, the packaging cell line disclosed in column 10, lines 4-7 of D21 must be interpreted as an implicit teaching of a cell line not comprising any pIX sequences. There is no reason for the person skilled in the art to interpret a "packaging cell line that comprises the sequences coding for E1A and the early proteins of the E1B region" to be a cell comprising the sequences coding for the late pIX protein.

(ii) The combination of the teaching in column 4 with the teaching in column 10

The OD states that there is allegedly no reason why the person skilled in the art would combine the teaching in column 4 relating to a vector with the teaching in column 10 relating to a packaging cell. The Appellant disagrees with this assertion.

According to the Guidelines for Examination, C-IV-9.9:

"In determining whether it would be obvious to combine two or more distinct disclosures, the examiner should also have regard in particular to the following:

(...)

(iii) the combining of two or more parts of the same document would be obvious if there is a reasonable basis for the skilled person to associate these parts with one another."

As will be shown below, in the present case there is "a reasonable basis" for the skilled person to associate the teaching in column 4 relating to a vector with the teaching in column 10 relating to a packaging cell.

Firstly, D21 relates to packaging systems for generating adenoviruses, wherein the system comprises an adenoviral vector *together with* a packaging cell in which the vector is used. This is clearly indicated in the first sentence of D21:

"The invention relates to new defective adenoviral vectors permitting the transfer and expression of genes of interest to a host eukaryotic cell or organism, as well as to new <u>cornplementation lines complementing in trans the essential viral functions which have been deleted from the genome of these recombinant adenoviruses."</u>
(column 1, lines 5-10 of D21a)

D21 thus relates to the combination of a defective adenoviral vector and of a packaging cell line providing in trans the functions which have been deleted in the corresponding adenoviral vector. Since the packaging cell line is designed to complement the vector, the person skilled in the art would not consider these two elements of the invention separately. Consequently, there is a basis in D21 to associate the teaching relating to a vector with the teaching relating to a packaging cell.

In addition, D21 emphasizes that the packaging cell line should not only be able to complement the vector, but should also be designed in such as way as to reduce or eliminate the adenoviral sequences in common with the vector:

"As regards an adenoviral vector according to the Invention derived from a human adenovirus type 5, said deletion comprises at least the sequences lying between nucleotides 1634 and 3509 of the adenoviral genome, the sequence of which is disclosed in the GenBank data bank under the reference M73260 [D3, editor's note]. The object of this deletion is to reduce or eliminate sequences which are common to an adenoviral vector according to the invention and the adenoviral genome fragment integrated in a complementation line, for example line 293." (column 4, lines 42-51 of D21a)

The above paragraph gives a clear motivation to the person skilled in the art to combine the position of the deletion present in the vector with the position of the adenoviral fragment present in the genome of the packaging cell in such a way as to reduce or eliminate sequences which are common to these two elements of the invention. As a matter of fact, during the oral proceedings, both the Patentee and the two Opponents agreed on the fact that the person skilled in the art would see the packaging cell as a "mirror image" of the adenoviral vector.

In view of the above, D21 not only provides a "reasonable basis" but provides a clear motivation for the skilled person to associate the teaching relating to a vector with the teaching relating to a packaging cell. In other worlds, the person skilled in the art immediately derived the packaging cell line from the vector.

(iii) The teaching in column 4 of D21a

The OD alleges that the following passage in column 4 of D21a teaches the skilled person to use a vector without pIX sequences in a cell comprising pIX sequences:

"As regards an adenoviral vector according to the invention derived from a human adenovirus type 5, said deletion comprises at least the sequences lying between nucleotides 1634 and 3509 of the adenoviral genome, the sequence of which is disclosed in the GenBank data bank under the reference M73260 [D3, editor's note]. The object of this deletion is to reduce or eliminate sequences which are common to an adenoviral vector according to the invention and the adenoviral genome fragment integrated in a complementation line, for example line 293." (column 4, lines 42-51)

The Appellant strongly disagrees with the OD's interpretation for the two reasons explained herebelow.

<u>Firstly</u>, the paragraph cited above teaches that the vector must comprise a deletion extending from nucleotide 1634 to nucleotide 3509 of Ad5. Since the pIX gene lies between nucleotides 3551 and 4070 of Ad5 (see e.g. D3), all pIX sequences are located *outside* of the deletion. Thus the above passage of D21 teaches the skilled person to use a vector comprising pIX sequences.

Since this paragraph further teaches that the sequences in common between the vector and the packaging line should be reduced or eliminated, the packaging cell line to be used with this vector should be devoid of pIX sequences. Thus the above paragraph teaches the skilled person to use a vector comprising pIX sequences in a complementation cell without pIX sequences, which is exactly the opposite of what is stated by the OD.

Secondly, the paragraph mentions that the cell line may for example be the 293 cell line, which is the only packaging cell line available in the prior art (see sentence bridging column 2 and column 3 of D21a). However, the statement that it is advantageous to reduce or eliminate common sequences is a general statement that is not limited to the 293 cell line. Specifically, D21 does not pertain to new adenoviral vectors to be used with the 293 cell line, but to new adenoviral vectors to be used with new packaging cell lines (see e.g. column 1, lines 5-10 of D21a). There is absolutely no reason to conclude that the general statement relating to overlapping sequences does not apply to the packaging cell lines of the invention.

(iv) Conclusion

In view of the above, it is submitted that the OD was wrong in concluding that the claims of AR2 are inventive.

4.1.3. Additional remarks

During the oral proceedings as well as in the written submissions, the Patentee tried to argue that the inventiveness of the claims stems from the absence of pIX sequences from the genome of the cell line. However, at the priority date of the opposed patent, the person skilled in the art knew that, while the cell must encode the E1A and the E1B early proteins, the presence of pIX within the cell is entirely optional.

For example, D53 (van den Elsen *et al.*, 1982) teaches a primary Baby Rat Kidney (BRK) cell co-transformed with the *Hpal-E* fragment (0-4.3%), which essentially comprises the E1A region, and with the *Smal-F* fragment (2.8-10.7%) which comprises most of the E1B region (see e.g. lines 3-8, page 177). As apparent from D3, the *Smal-F* fragment comprising most of the E1B region corresponds to nucleotides 1045-3978 of Ad5. Since the sequence coding for pIX extends from nucleotides 3609 to 4028 of Ad5, the cell line of D53 lacks 50 nucleotides at the 3' end of the pIX coding sequence. The cell of D53 thus lacks pIX sequences, and only codes for a truncated pIX protein.

D54 (Kawarabayasi and Sugisaki, 1985) also teaches a primary cell comprising an adenovirus fragment that lacks pIX sequences. D54 teaches a primary rat cell transformed with a fragment consisting of Ad12 nucleotides 118-3520 (see e.g. abstract, line 4-5). As evidenced by D55, the sequence coding for pIX extends from nucleotides 3374 to 3808 of Ad12 (see e.g. Figure 3 on pages 6579-6580 of D55). Thus 288 nucleotides of the 3' end of the pIX coding sequence (i.e. 66%) are lacking in this cell line. Therefore, the cell of D53 comprises in its genome an Ad12 fragment encoding the E1A and the E1B early proteins, but only a truncated, non-functional portion of the pIX protein.

In addition, D46 teaches that pIX has to be present in the replicating vector genome to be properly expressed (see e.g. the abstract of D46). As a consequence, if one wishes to eliminate overlaps between vector DNA and adenovirus DNA inserted into the genome, pIX must be present in the viral vector and not in the cell genome.

In view of the prior art at the priority date of the opposed patent, there was therefore nothing original or inventive with the provision of a primary cell that encodes the E1 functions but not the pIX protein.

4.2. Claims 1-16 and 18 of AR2

Independent claims 1 is directed to a combination of the packaging cell of claim 19 with an adenoviral vector, wherein said adenoviral vector has no overlapping sequences which allow for homologous recombination between the cell and the vector. Since this claim is based on the packaging cell of claim 19, and since D21 teaches to eliminate the sequences in common between the cell and the vector (see e.g. column 4, lines 42-51 of D21a), it must be concluded that its subject matter does not involve an inventive step.

Claims 2, 3, 5-9, 11-13, 15 and 18 all clepend either directly or indirectly on claim 1. The additional features present in these claims do not add subject-matter that goes beyond the knowledge of the person skilled in the art. Therefore, these claims lack inventiveness as well.

Claim 4 is directed to a combination of a packaging cell with an adenoviral vector, wherein said combination comprises an adenoviral vector resulting from the action of a nucleic acid polymerase comprised by the combination of claim 3. This claims is contrary to Art. 56 EPC for the same reasons as claim 3.

Claim 10 is redundant with claim 19, so the arguments presented above under section 4.1 apply *mutatis mutandis* to this claim.

Claim 16 is directed to a specific cell according to claim 10 that has been deposited at the ECACC. This cell has no additional feature that would render it inventive.

In view of the above, claims 1-16 and 18 of AR2 do not fulfill the requirements of Art. 100(a) and 56 EPC.

5. Art. 100(c) and Art. 100(b) EPC

The Patentee has filed an appeal against the OD's decision to the extend it was adverse to the patent proprietor. Insofar as the Patentee defends its patent in a scope broader than the patent as maintained, it is submitted that the patent as granted is contrary to Art. 100(c) and 100(b) EPC.

In order to avoid any unnecessary repetition, we refer to our submissions filed during the opposition, including all cited documents, demonstrating why the patent in a scope broader than the patent as maintained does not comply with Art. 100(c) and 100(b) EPC.

6. Conclusion

A packaging cell comprising in its genome an adenoviral sequence consisting of Ad5 nucleotides 459-3510 lacks inventiveness, and therefore all claims but claim 17 of the opposed patent as maintained in amended form are contrary to Art. 100(a) and 56 EPC.

In addition, claim 17 of the opposed patent as maintained in amended form is contrary to Art. 100(a) and 54 EPC, as shown in section 3 of the present submission.

Therefore, the request for revocation of the opposed patent in its entirety is justified.

Please stamp and return the enclosed EPO Form 1037 as evidence of safe receipt of this letter and its enclosures.

Yours truly,

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Chrote Peller

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- Encl.: Figure 2 by Opponent 01 (D52);
 - Van den Elsen et al., 1982 (D53);
 - Kawarabayasi and Sugisaki, 1985 (D54);
 - Kimura et al., 1981 (D55);
 - EPO forms 1037.1 and 1037.2.